

NUS CORPORATION
SUPERFUND DIVISION

10816
FILE COPY

INTERNAL CORRESPONDENCE

C-583-7-4-10

TO: ED TAYLOR

DATE: JULY 5, 1984

FROM: HANS-PETER KRAHN *HK*

COPIES: FILE

SUBJECT: CAMBRIDGE ANALYTICAL DATA VALIDATION FOR INTERSTATE/WOBURN

Samples: A 2749 - A 2757

Case No. 2660

TDD No. F1-8303-07A

Job No. 3425-03

0300.01

Superfund Records Center

SITE: Wells, E+H

BREAK: 8.2.3

OTHER: 531618



SDMS DocID

531618

The organic analytical data from Cambridge Analytical Associates (Case 2660) has been reviewed in accordance with Region I FIT Standard Operating Procedure No. 4 to insure the validity of the data presented. This evaluation for samples A 2749 through A2757 was based on Level I analysis considering the following parameters:

- data completeness
- laboratory and field blanks
- laboratory and field duplicates
- surrogate spikes
- matrix spikes
- spectral performance
- holding times

The data package contained all the pertinent information requested to accurately assess the accuracy and precision of the samples sent for analysis with only one exception: standard reference spectra was not provided. The sample spectra was therefore matched by the reviewer with the same compounds from the EPA/NIH Mass Spectral Data Base where the match was found to be satisfactory. No contaminants were found to be above the contractual detection limits in either the reagent or blind EPA lab blank. Methylene chloride and acetone were found in trace levels, and toluene also was present at a low concentration in the blind blank. The blind field blank was contaminated with chloroform and acetone and, therefore, those compounds should be rejected. Methylene chloride, toluene, and trichloroethylene were found in trace quantities below the method detection limits and are not found at higher levels in the samples so no action should be taken. The laboratory sample precision was of good quality, with no duplicate results exceeding the contractual criteria. The field duplicates were too low to be meaningful. A few of the surrogate spikes for toluene-d₈ were outside the acceptable range, but the recoveries were close and would not justify taking any action. The matrix spikes all had very good recoveries. Holding times were also met.

C-583-7-4-10

MEMO TO: ED TAYLOR
JULY 5, 1984 - PAGE TWO

Listed below is a summary of the recommendations for this case (No. 2660):

<u>Compound</u>	<u>Action</u>
acetone	R
chloroform	R

R = Rejected

I hereby acknowledge receipt and approval of the NUS/FIT Level I data validation for TDD F1-8303-07A (Case 2660) from Cambridge Analytical Associates.


Deputy Project Officer
Regional Sample Control Center

HPK/tao

cc: D. Smith
R. DiNitto
J. Morin
A. DeMarco

500 m

NUS CORPORATION
SUPERFUND DIVISION

C-583-74-10 PROJECT NOTES

Q/A 7/3/84

TO: Ed Taylor DATE: 28 June 1984

FROM: H-P Krahn COPIES: File

SUBJECT: Cambridge Analytical
~~Interstate~~ Data Validation for Interstate/Woburn

REFERENCE: Case # 2660

TDD # F1-8303-07A

JOB # 3425-03

The organic analytical data from Cambridge Analytical Associates (Case 2660) has been reviewed ^{in accordance with Region I SFF standard operating procedure #4} to insure the validity of the data presented.

This evaluation for samples A2749 through A2757 was based on Level I analysis considering the following parameters:

- data completeness
- laboratory + field blanks
- labs + field duplicates
- surrogate spikes
- matrix spikes
- spectral performance
- holding times

The data package contained all the pertinent information requested to accurately assess the accuracy + precision of the samples sent for analysis with only one exception: standard reference spectra was not provided

The sample spectra was ^{therefore} ~~matched~~ ^{by the ~~lower~~ with} the same compounds from the EPA/NIH Mass Spectral Data Base where the match was found to be satisfactory.

~~The laboratory blanks were found to be free of all contaminants and the~~ No contaminants were found to be above ^{blind} the contractual detection limits in either the reagent or EPA lab blanks.

Methylene Chloride and acetone were found in trace levels, and toluene also was present ^{at a} ~~is~~ very low concentration in the blind blank.

The ^{blind} field blank was ~~however~~ contaminated with chloroform

R?
A?

and acetone, ^{+ therefore} ~~concluding that~~ those compounds ^{should} to be rejected. Methylene chloride, toluene, and trichloroethene were found in trace quantities below the laboratory sample precision was of good quality, ^{with} ~~no~~ duplicates ~~being~~ ^{results} exceeding the contractual criteria. The field duplicates were too low to be meaningful. ~~A few of the~~ ^{some} ~~matrix~~ ^{matrix} spikes for Toluene-d₈ were outside the acceptable range, but the recoveries were close and would not justify ~~any~~ taking any action. The matrix spikes all had very good recoveries. Holding times were also met.

Method detection limits so no action should be taken.
 Some are not found at higher levels in samples.

Listed below is a summary of the recommendations for this case (No. 2660):

<u>Compound</u>	<u>Action</u>
acetone	R
chloroform	R

R = rejected

I hereby acknowledge receipt and approval of the NUS/FIT Level 1 data validation for TDD FI-8303-07A (Case 2660) from Cambridge Analytical Associates.

Deputy Project Officer
Regional Sample Control Center

cc: D. Smith
R. DeBrito
J. Marin
A. Delanceo



Cambridge Analytical Associates

222 Arsenal Street / Watertown, Massachusetts 02172 / (617)923-9376

F.I.L.

June 1, 1984

*Interstate
Joanne Morin.*

Ms. Leslie Braun
US Environmental Protection Agency
CLP Sample Management Office
VIAR & Co.
300 Lee Street
Alexandria, VA 22314

Dear Leslie:

Enclosed please find the results of our analysis of samples received as case number 2660. These analyses were performed as part of our contract number 68-01-6791.

The samples were analyzed for volatile organics only. There were no unusual problems encountered during the analysis.

Please note that we elected not to screen the samples by GC/FID prior to analysis.

If you should have any questions, please feel free to call.

Sincerely,

A handwritten signature in cursive script that reads "Scott R. Drew".

Scott R. Drew
Senior Scientist

Enclosure
mjk

DELIVERABLES INDEX

CASE # 2660

CONTRACTOR CAMBRIDGE ANALYTICAL ASSOCIATES CONTRACT # 68-01-6791

NOTE: VOLATILE ORGANICS ONLY

SECTION TITLE	
I	<p><u>NARRATIVE</u></p> <p>Contains: Case #, Contract #, summary of any QC, sample, shipment and analytical problems, documentation of any internal decision tree process used.</p>
II	<p><u>QC SUMMARY</u></p> <p>*A. Surrogate % recovery summary form.</p> <p>*B. Reagent blank summary form.</p> <p>*C. Matrix spike duplicate/recovery form</p> <p>*D. Instrument tune & performance summary form</p>
III	<p><u>SAMPLE DATA PACKET</u></p> <p>A. <u>Sample Data in increasing SMO # order:</u></p> <ol style="list-style-type: none"> 1) HSL results. 2) GC/MS tentative ID sheet, even if none found. 3) Raw data (VOA, A/B/N, Pest, Dioxin) <ol style="list-style-type: none"> i. Chromatogram(s) ii. Data System printout iii. HSL spectra with standard (dual display) iv. GC/MS library search spectra v. Quantitation/calculation of tentative ID concentration
IV	<p><u>STANDARDS PACKET</u></p> <ol style="list-style-type: none"> A. Cross-reference table (lab's internal). B. VOA standards chromatograms and data system printouts. C. A/B/N standards chromatograms and data system printouts. D. Pesticide standards chromatograms and data system printouts. E. Dioxin standards chromatograms and data system printouts. *F. Internal standard verification data sheet. *G. FSGG initial calibration data. *H. Calibration check. I. Current list of lab detection limits.

*Designates new form.

cont.

SECTION TITLE

V

RAW QC DATA PACKET

A. DFTPP

- ~~*1) GC/MS performance standard form.~~
- ~~2) Bar graph spectrum.~~
- ~~3) M/Z listing.~~

B. BFB

- *1) GC/MS performance standard form.
- 2) Bar graph spectrum.
- 3) M/Z listing.

C. Chromatography Check Data

- 1) Benzidine EICP's.
- 2) Pentachlorophenol EICP's/calculations.
- 3) GC/EC column check.

D. Blank Data(in order VOA, A/B/N, Pesticide, Dioxin)

- 1) Chromatograms.
- 2) Data system printout.

E. Matrix Spike Data(in order VOA, A/B/N, Pesticide)

- 1) Chromatograms.
- 2) Data system printout.

F. Matrix Spike Duplicate Data(in order VOA, A/B/N, Pesticide)

- 1) Chromatograms.
- 2) Data system printout.

VI

SAMPLE PREPARATION PACKET

- A. Sample control sheet (lab's internal).
- B. Screen data(in order VOA, A/B/N, pesticide)

- 1) GC/FID chromatograms.
- 2) GC screen data sheets.

*Designates new form.

00002

REAGENT BLANK SUMMARY

CASE NO. 2660
 LOW LEVEL X
 WATER X
 QC REPORT NO. 011

CONTRACTOR Cambridge Analytical
 MED. LEVEL _____
 SOIL/SED. _____

CONTRACT NO. 68-01-6791
 HIGH LEVEL _____
 OTHER (Specify) _____
 UNITS (Circle) ug/kg ug/l

FRACTION	CAS NUMBER	COMPOUND	CONCENTRATION	CONTRACT DETECTION LIMITS	COMMENTS
VOLATILES:	<u>75-09-2</u>	<u>Methylene Chloride</u>	<u>2.3</u>	<u>5</u>	<u>Airborne or MeOH</u>
	<u>67-64-1</u>	<u>Acetone</u>	<u>3.1</u>	<u>5</u>	<u>Airborne or MeOH</u>
File I.D. <u>CLPVOA 32</u>					
Instrument I.D. <u>OWA</u>					
SEMI-VOLATILES:					
File I.D.					
Instrument I.D.					
PESTICIDES:					
File I.D.					
Instrument I.D.					

FORM IV

00004

ea

MATRIX SPIKE DUPLICATE/RECOVERY

CASE NO. 2660
 LOW LEVEL X
 WATER X
 QC REPORT NO. 011

CONTRACTOR Cambridge Analytical
 MED. LEVEL _____
 SOIL/SED. _____

CONTRACT NO. 68-01-6791
 HIGH LEVEL _____
 OTHER (Specify) _____
 UNITS (Circle) ug/l

FRACTION	COMPOUND	CONC. SPTKE ADDED	CONC. MS	% REC.	CONC. MSD	% REC.	RPD	QC LIMITS*		COMMENTS
								RPD	RECOVERY	
VOA SMD # A2749	1,1-Dichloroethylene	25	11.9	48*	13.2	53	10.3	<15%	51-151	
	Trichloroethylene	25	21.9	88	22.4	90	2.3	<15%	74-128	
	Chlorobenzene	25	22.7	91	26.6	106	15.8*	<15%	67-131	
	Toluene	25	25.1	100	26.6	106	5.8	<15%	58-132	
	Benzene	25	25.0	100	26.3	105	5.1	<15%	56-132	
B/N SMD #	1,2,4-Trichlorobenzene							<50%	38-108	
	Acenaphthene							<50%	57-115	
	2,4-Dinitrotoluene							<50%	43-113	
	Di-N-Butylphthalate							<50%	13-113	
	Pyrene							<50%	25-137	
	N-Nitrosodi-N-Propylamine							<50%	34-114	
ACID SMD #	1,4-Dichlorobenzene							<40%	33-103	
	Pentachlorophenol							<40%	19-123	
	Phenol							<40%	23-81	
	2-Chlorophenol							<40%	33-107	
	p-Chloro-m-Cresol							<40%	32-108	
PEST SMD #	4-Nitrophenol							<40%	15-93	
	Lindane							<40%	87-107	
	Heptachlor							<40%	43-125	
	Aldrin							<40%	45-109	
	Dieldrin							<40%	56-122	
	Endrin							<40%	89-101	
	p,p-DDT							<40%	82-102	

*Asterisked values are outside QC limits.

RPD: VOAs 1 out of 5; outside QC limits
 B/N _____ out of _____; outside QC limits
 ACID _____ out of _____; outside QC limits
 PEST _____ out of _____; outside QC limits

RECOVERY: VOAs 1 out of 10; outside QC limits
 B/N _____ out of _____; outside QC limits
 ACID _____ out of _____; outside QC limits
 PEST _____ out of _____; outside QC limits

*Date Limits Set 12/82
 Revision Due 6/83

FORM V

00005



Cambridge Analytical Associates

SAMPLE DATA PACKET

00006

A2749

ORGANICS ANALYSIS DATA SHEET

Laboratory Name: Cambridge Analytical Associates
Lab Sample ID No: 8402-434
Sample Matrix: Water
Data Release Authorized By: [Signature]

Case No: 2660
QC Report No: 011
Contract No.: 68-01-6791
Date Sample Received: 5/3/84

VOLATILES

CONCENTRATION: LOW MEDIUM HIGH (circle one)
DATE EXTRACTED/PREPARED:
DATE ANALYZED: 5/5/84
PERCENT MOISTURE: 100
CONC./DILUTION FACTOR: 5:1

PESTICIDES

CONCENTRATION: LOW MEDIUM HIGH (circle one)
DATE EXTRACTED/PREPARED:
DATE ANALYZED:
PERCENT MOISTURE:
CONC./DILUTION FACTOR:

NOT ANALYZED

Table with columns PP #, CAS #, and chemical name. Includes handwritten notes like 'ET 3/24/84' and '100U' for various compounds such as acrolein, benzene, and chloroform.

Table with columns PP #, CAS #, and chemical name. Lists pesticides like aldrin, dieldrin, and various BHCs, all marked as 'NOT ANALYZED'.

DIOXINS

CONCENTRATION: LOW MEDIUM HIGH (circle one)
DATE EXTRACTED/PREPARED:
DATE ANALYZED:
PERCENT MOISTURE:
CONC./DILUTION FACTOR:

Table with columns PP #, CAS #, and chemical name. Includes entry (129B) 1746-01-6 2,3,7,8-tetrachlorodibenzo-p-dioxin.

Sample Number
A2749

Laboratory Name: Cambridge Analytical Case No: 2660
 QC Report No: 011

B. Tentatively Identified Compounds

CAS #	Compound Name	Fraction	Scan No. or Retention Time	% Maximum Score Attained Mass Matching Routine: (Specify: _____)	Estimated Concentration (ug/L or ug/kg)
1.	NONE DETECTED	VOA			
2.					
3.					
4.					
5.					
6.					
7.					
8.					
9.					
10.					
11.					
12.					
13.					
14.					
15.					
16.					
17.					
18.					
19.					
20.					
21.					
22.					
23.					
24.					
25.					
26.					
27.					
28.					
29.					
30.					

6/22

A2750

ORGANICS ANALYSIS DATA SHEET

Laboratory Name: Cambridge Analytical Associates
 Lab Sample ID No: 8402-435
 Sample Matrix: Water
 Data Release Authorized By: [Signature]

Case No: 2660
 QC Report No: 011
 Contract No.: 68-01-6791
 Date Sample Received: 5/3/84

VOLATILES

CONCENTRATION: (LOW) MEDIUM HIGH (circle one)
 DATE EXTRACTED/PREPARED: _____
 DATE ANALYZED: 5/5/84
 PERCENT MOISTURE: 100
 CONC./DILUTION FACTOR: 5:1

PESTICIDES

CONCENTRATION: LOW MEDIUM HIGH (circle one)
 DATE EXTRACTED/PREPARED: _____
 DATE ANALYZED: _____
 PERCENT MOISTURE: _____
 CONC./DILUTION FACTOR: _____

NOT ANALYZED

PP #	CAS #	Chemical Name	Concentration (ug/l or ug/kg, circle one)
(2V)	107-02-8	acrolein	100U
(3V)	107-13-1	acrylonitrile	100U
(4V)	71-43-2	benzene	SU
(6V)	56-23-5	carbon tetrachloride	SU
(7V)	108-90-7	chlorobenzene	SU
(10V)	107-06-2	1,2-dichloroethane	1U
(11V)	71-55-6	1,1,1-trichloroethane	SU
(13V)	75-34-3	1,1-dichloroethane	SU
(14V)	79-00-5	1,1,2-trichloroethane	SU
(15V)	79-34-5	1,1,2,2-tetrachloroethane	10U
(16V)	75-00-3	chloroethane	10U
(19V)	110-75-8	2-chloroethylvinyl ether	10U
(23V)	67-66-3	chloroform	SU R
(29V)	75-35-4	1,1-dichloroethene	SU
(30V)	156-60-5	trans-1,2-dichloroethene	SU
(32V)	78-87-5	1,2-dichloropropane	10U
(33V)	10061-02-6	trans-1,3-dichloropropene	SU
	10061-01-05	cis-1,3-dichloropropene	SU
(38V)	100-41-4	ethylbenzene	SU
(44V)	75-09-2	methylene chloride	El 50 SK
(45V)	74-87-3	chloromethane	10U
(46V)	74-83-9	bromomethane	10U
(47V)	75-25-2	bromoform	10U
(48V)	75-27-4	bromodichloromethane	SU
(49V)	75-69-4	fluorotrichloromethane	---
(50V)	75-71-8	dichlorodifluoromethane	---
(51V)	124-48-1	chlorodibromomethane	SU
(55V)	127-18-4	tetrachloroethene	SU
(86V)	108-88-3	toluene	SU
(87V)	79-01-6	trichloroethene	SU
(88V)	75-01-4	vinyl chloride	10U
	67-64-1	acetone	El 50 SK R
	78-93-3	2-butanone	SU
	75-15-0	carbonyl disulfide	1U
	519-78-6	2-hexanone	SU
	108-10-1	4-methyl-2-pentanone	SU
	100-42-5	styrene	SU
	108-05-4	vinyl acetate	SU
	1330-20-7	total xylenes	SU

PP #	CAS #	Chemical Name	Concentration (ug/l or ug/kg, circle one)
(89P)	309-00-2	aldrin	
(90P)	60-57-1	dieldrin	
(91P)	57-74-9	chlordane	
(92P)	50-29-3	4,4'-DDT	
(93P)	72-55-9	4,4'-DDE	
(94P)	72-54-8	4,4'-DDD	
(95P)	115-29-7	α-endosulfan	
(96P)	115-29-7	β-endosulfan	
(97P)	1031-07-8	endosulfan sulfate	
(98P)	72-20-8	endrin	
(99P)	7421-93-4	endrin aldehyde	
(100P)	76-44-8	heptachlor	
(101P)	1024-57-3	heptachlor epoxide	
(102P)	319-84-6	α-BHC	
(103P)	319-85-7	β-BHC	
(104P)	319-86-8	δ-BHC	
(105P)	58-89-9	γ-BHC (lindane)	
(106P)	53469-21-9	PCB-1242	
(107P)	11097-69-1	PCB-1254	
(108P)	11104-28-2	PCB-1221	
(109P)	11141-16-5	PCB-1232	
(110P)	12672-29-6	PCB-1248	
(111P)	11096-82-5	PCB-1260	
(112P)	12674-11-2	PCB-1016	
(113P)	8001-35-2	toxaphene	

DIOXINS

CONCENTRATION: LOW MEDIUM HIGH (circle one)
 DATE EXTRACTED/PREPARED: _____
 DATE ANALYZED: _____
 PERCENT MOISTURE: _____
 CONC./DILUTION FACTOR: _____

PP #	CAS #	Chemical Name	Concentration (ug/l or ug/kg, circle one)
(129B)	1746-01-6	2,3,7,8-tetrachlorodibenzo-p-dioxin	

00009

Sample Number
A2750

Laboratory Name: Cambridge Analytical Case No: 2660
 QC Report No: 011

B. Tentatively Identified Compounds

CAS #	Compound Name	Fraction	Scan No. or Retention Time	% Maximum Score Attained Mass Matching Routine: (Specify: _____)	Estimated Concentration (ug/L or ug/g)
1.	NONE DETECTED	NOA			
2.					
3.					
4.					
5.					
6.					
7.					
8.					
9.					
10.					
11.					
12.					
13.					
14.					
15.					
16.					
17.					
18.					
19.					
20.					
21.					
22.					
23.					
24.					
25.					
26.					
27.					
28.					
29.					
30.					

•/12

A 2751

ORGANICS ANALYSIS DATA SHEET

Laboratory Name: Cambridge Analytical Associates
 Lab Sample ID No: 8402-436
 Sample Matrix: Water
 Data Release Authorized By: [Signature]

Case No: 2660
 QC Report No: 011
 Contract No.: 68-01-6791
 Date Sample Received: 5/3/84

VOLATILES

CONCENTRATION: (LOW) MEDIUM HIGH (circle one)
 DATE EXTRACTED/PREPARED: _____
 DATE ANALYZED: 5/5/84
 PERCENT MOISTURE: 100
 CONC./DILUTION FACTOR: 5:1

EL 2/24/84 ug/l
 or ug/kg
 (circle one)
100U

PP #	CAS #	Compound	Concentration
(2V)	107-02-8	acrolein	
(3V)	107-13-1	acrylonitrile	100U
(4V)	71-43-2	benzene	SU
(6V)	56-23-5	carbon tetrachloride	SU
(7V)	108-90-7	chlorobenzene	SU
(10V)	107-06-2	1,2-dichloroethane	1U
(11V)	71-55-6	1,1,1-trichloroethane	SU
(13V)	75-34-3	1,1-dichloroethane	SU
(14V)	79-00-5	1,1,2-trichloroethane	SU
(15V)	79-34-5	1,1,2,2-tetrachloroethane	10U
(16V)	75-00-3	chloroethane	10U
(19V)	110-75-8	2-chloroethylvinyl ether	10U
(23V)	67-66-3	chloroform	SU R
(29V)	75-35-4	1,1-dichloroethene	SU
(30V)	156-60-5	trans-1,2-dichloroethene	SU
(32V)	78-87-5	1,2-dichloropropane	10U
(33V)	10061-02-6	trans-1,3-dichloropropene	SU
	10061-01-05	cis-1,3-dichloropropene	SU
(38V)	100-41-4	ethylbenzene	SU
(44V)	75-09-2	methylene chloride	EL SU SK
(45V)	74-87-3	chloromethane	10U
(46V)	74-83-9	bromomethane	10U
(47V)	75-25-2	bromoform	10U
(48V)	75-27-4	bromodichloromethane	SU
(49V)	75-69-4	fluorotrichloromethane	---
(50V)	75-71-8	dichlorodifluoromethane	---
(51V)	124-48-1	chlorodibromomethane	SU
(85V)	127-18-4	tetrachloroethene	SU
(86V)	108-88-3	toluene	SU
(87V)	79-01-6	trichloroethene	SU
(88V)	75-01-4	vinyl chloride	10U
	67-64-1	acetone	EL SU SK
	78-93-3	2-butanone	SU
	75-15-0	carbendisulfide	1U
	519-78-6	2-hexanone	SU
	108-10-1	4-methyl-2-pentanone	SU
	100-42-5	styrene	SU
	108-05-4	vinyl acetate	SU
	1330-20-7	total xylenes	SU

PESTICIDES

CONCENTRATION: LOW MEDIUM HIGH (circle one)
 DATE EXTRACTED/PREPARED: _____
 DATE ANALYZED: _____
 PERCENT MOISTURE: _____
 CONC./DILUTION FACTOR: _____

NOT ANALYZED

PP #	CAS #	Compound	Concentration
(89P)	309-00-2	aldrin	
(90P)	60-57-1	dieldrin	
(91P)	57-74-9	chlordan	
(92P)	50-29-3	4,4'-DDT	
(93P)	72-55-9	4,4'-DDE	
(94P)	72-54-8	4,4'-DDD	
(95P)	115-29-7	α -endosulfan	
(96P)	115-29-7	β -endosulfan	
(97P)	1031-07-8	endosulfan sulfate	
(98P)	72-20-8	endrin	
(99P)	7421-93-4	endrin aldehyde	
(100P)	76-44-8	heptachlor	
(101P)	1024-57-3	heptachlor epoxide	
(102P)	319-84-6	α -BHC	
(103P)	319-85-7	β -BHC	
(104P)	319-86-8	δ -BHC	
(105P)	58-89-9	γ -BHC (lindane)	
(106P)	53469-21-9	PCB-1242	
(107P)	11097-69-1	PCB-1254	
(108P)	11104-28-2	PCB-1221	
(109P)	11141-16-5	PCB-1232	
(110P)	12672-29-6	PCB-1248	
(111P)	11096-82-5	PCB-1260	
(112P)	12674-11-2	PCB-1016	
(113P)	8001-35-2	toxaphene	

DIOXINS

CONCENTRATION: LOW MEDIUM HIGH (circle one)
 DATE EXTRACTED/PREPARED: _____
 DATE ANALYZED: _____
 PERCENT MOISTURE: _____
 CONC./DILUTION FACTOR: _____

PP #	CAS #	Compound	Concentration
(129B)	1746-01-6	2,3,7,8-tetrachlorodibenzo-p-dioxin	

00011

Sample Number
A2751

Laboratory Name: Cambridge Analytical Case No: 2660
 QC Report No: 011

B. Tentatively Identified Compounds

CAS #	Compound Name	Fraction	Scan No. or Retention Time	% Maximum Score Attained Mass Matching Routine: (Specify: _____)	Estimated Concentration (ug/L or ug/g)
1.	NONE DETECTED	NDA			
2.					
3.					
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30.					

6/22

2752

ORGANICS ANALYSIS DATA SHEET

Laboratory Name: Cambridge Analytical Associates
 Lab Sample ID No: 8402-437
 Sample Matrix: Water
 Data Release Authorized By: [Signature]

Case No: 2660
 QC Report No: 011
 Contract No.: 68-01-6791
 Date Sample Received: 5/3/84

VOLATILES

CONCENTRATION: (LOW) MEDIUM HIGH (circle one)
 DATE EXTRACTED/PREPARED: _____
 DATE ANALYZED: 5/5/84
 PERCENT MOISTURE: 100
 CONC./DILUTION FACTOR: 5:1

PESTICIDES

CONCENTRATION: LOW MEDIUM HIGH (circle one)
 DATE EXTRACTED/PREPARED: _____
 DATE ANALYZED: _____
 PERCENT MOISTURE: _____
 CONC./DILUTION FACTOR: _____

PP #	CAS #	Compound	ug/l or ug/kg (circle one)
(2V)	107-02-8	acrolein	100U
(3V)	107-13-1	acrylonitrile	100U
(4V)	71-43-2	benzene	SU
(6V)	56-23-5	carbon tetrachloride	SU
(7V)	108-90-7	chlorobenzene	SU
(10V)	107-06-2	1,2-dichloroethane	1U
(11V)	71-55-6	1,1,1-trichloroethane	SU
(13V)	75-34-3	1,1-dichloroethane	SU
(14V)	79-00-5	1,1,2-trichloroethane	SU
(15V)	79-34-5	1,1,2,2-tetrachloroethane	10U
(16V)	75-00-3	chloroethane	10U
(19V)	110-75-8	2-chloroethylvinyl ether	10U
(23V)	67-66-3	chloroform	SUR
(29V)	75-35-4	1,1-dichloroethene	SU
(30V)	156-60-5	trans-1,2-dichloroethene	SU
(32V)	78-87-5	1,2-dichloropropane	10U
(33V)	10061-02-6	trans-1,3-dichloropropene	SU
	10061-01-05	cis-1,3-dichloropropene	SU
(38V)	100-41-4	ethylbenzene	SU
(44V)	75-09-2	methylene chloride	SU SK
(45V)	74-87-3	chloromethane	10U
(46V)	74-83-9	bromomethane	10U
(47V)	75-25-2	bromoform	10U
(48V)	75-27-4	bromodichloromethane	SU
(49V)	75-69-4	fluorotrichloromethane	---
(50V)	75-71-8	dichlorodifluoromethane	---
(51V)	124-48-1	chlorodibromomethane	SU
(83V)	127-18-4	tetrachloroethene	SU
(86V)	108-88-3	toluene	SU SK
(87V)	79-01-6	trichloroethene	SU
(88V)	75-01-4	vinyl chloride	10U
	67-64-1	acetone	SU 8BR
	78-93-3	2-butanone	SU
	75-15-0	carbendisulfide	1U
	519-78-6	2-hexanone	SU
	108-10-1	4-methyl-2-pentanone	SU
	100-42-5	styrene	SU
	108-05-4	vinyl acetate	SU
	1330-20-7	total xylenes	SU

E1 7/21/84 (circle one) ug/l or ug/kg

NOT ANALYZED

PP #	CAS #	Compound	ug/l or ug/kg (circle one)
(89P)	309-00-2	aldrin	
(90P)	60-57-1	dieldrin	
(91P)	57-74-9	chlordan	
(92P)	50-29-3	4,4'-DDT	
(93P)	72-55-9	4,4'-DDE	
(94P)	72-54-8	4,4'-DDD	
(95P)	115-29-7	α -endosulfan	
(96P)	115-29-7	β -endosulfan	
(97P)	1031-07-8	endosulfan sulfate	
(98P)	72-20-8	endrin	
(99P)	7421-93-4	endrin aldehyde	
(100P)	76-44-8	heptachlor	
(101P)	1024-57-3	heptachlor epoxide	
(102P)	319-84-6	α -BHC	
(103P)	319-85-7	β -BHC	
(104P)	319-86-8	δ -BHC	
(105P)	58-89-9	γ -BHC (lindane)	
(106P)	53469-21-9	PCB-1242	
(107P)	11097-69-1	PCB-1254	
(108P)	11104-28-2	PCB-1221	
(109P)	11141-16-5	PCB-1232	
(110P)	12672-29-6	PCB-1248	
(111P)	11096-82-5	PCB-1260	
(112P)	12674-11-2	PCB-1016	
(113P)	8001-35-2	toxaphene	

DIOXINS

CONCENTRATION: LOW MEDIUM HIGH (circle one)
 DATE EXTRACTED/PREPARED: _____
 DATE ANALYZED: _____
 PERCENT MOISTURE: _____
 CONC./DILUTION FACTOR: _____

PP #	CAS #	Compound	ug/l or ug/kg (circle one)
(129B)	1746-01-6	2,3,7,8-tetrachlorodibenzo-p-dioxin	

Sample Number
A2752

Laboratory Name: Cambridge Analytical Case No: 2660
 QC Report No: 011

B. Tentatively Identified Compounds

CAS #	Compound Name	Fraction	Scan No. or Retention Time	% Maximum Score Attained Mass Matching Routine: (Specify: <u>PURITY</u>)	Estimated Concentration (ug/L or ug/kg)
1.	<u>3-Methoxy-3-Methyl-2-Butanone</u>	<u>VDA</u>	<u>811</u>	<u>91.4 %</u>	<u>5</u>
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A2753

ORGANICS ANALYSIS DATA SHEET

Laboratory Name: Cambridge Analytical Associates
 Lab Sample ID No: 8402-438
 Sample Matrix: Water
 Data Release Authorized By: [Signature]

Case No: 2660
 QC Report No: 011
 Contract No.: 68-01-6791
 Date Sample Received: 5/3/84

VOLATILES

CONCENTRATION: (LOW) MEDIUM HIGH (circle one)
 DATE EXTRACTED/PREPARED: _____
 DATE ANALYZED: 5/5/84
 PERCENT MOISTURE: 100
 CONC./DILUTION FACTOR: 5:1

E1
7/21/84 ug/l
 or ug/kg
 (circle one)
100U

PP #	CAS #	NAME	CONC.
(2V)	107-02-8	acrolein	
(3V)	107-13-1	acrylonitrile	<u>100U</u>
(4V)	71-43-2	benzene	<u>SU</u>
(6V)	56-23-5	carbon tetrachloride	<u>SU</u>
(7V)	108-90-7	chlorobenzene	<u>SU</u>
(10V)	107-06-2	1,2-dichloroethane	<u>1U</u>
(11V)	71-55-6	1,1,1-trichloroethane	<u>SU</u>
(13V)	75-34-3	1,1-dichloroethane	<u>SU</u>
(14V)	79-00-5	1,1,2-trichloroethane	<u>SU</u>
(15V)	79-34-5	1,1,2,2-tetrachloroethane	<u>10U</u>
(16V)	75-00-3	chloroethane	<u>10U</u>
(19V)	110-75-8	2-chloroethylvinyl ether	<u>10U</u>
(23V)	67-66-3	chloroform	<u>SUR</u>
(29V)	75-35-4	1,1-dichloroethene	<u>SU</u>
(30V)	156-60-5	trans-1,2-dichloroethene	<u>SU</u>
(32V)	78-87-5	1,2-dichloropropane	<u>10U</u>
(33V)	10061-02-6	trans-1,3-dichloropropene	<u>SU</u>
	10061-01-05	cis-1,3-dichloropropene	<u>SU</u>
(38V)	100-41-4	ethylbenzene	<u>SU</u>
(44V)	75-09-2	methylene chloride	<u>50-5K</u>
(45V)	74-87-3	chloromethane	<u>10U</u>
(46V)	74-83-9	bromomethane	<u>10U</u>
(47V)	75-25-2	bromoform	<u>10U</u>
(48V)	75-27-4	bromodichloromethane	<u>SU</u>
(49V)	75-69-4	fluorotrichloromethane	_____
(50V)	75-71-8	dichlorodifluoromethane	_____
(51V)	124-48-1	chlorodibromomethane	<u>SU</u>
(55V)	127-18-4	tetrachloroethene	<u>SU</u>
(86V)	108-88-3	toluene	<u>SU</u>
(87V)	79-01-6	trichloroethene	<u>SU</u>
(88V)	75-01-4	vinyl chloride	<u>10U</u>
	67-64-1	acetone	<u>50-5K</u>
	78-93-3	2-butanone	<u>SU</u>
	75-15-0	carbonylsulfide	<u>1U</u>
	519-78-6	2-hexanone	<u>SU</u>
	108-10-1	4-methyl-2-pentanone	<u>SU</u>
	100-42-5	styrene	<u>SU</u>
	108-05-4	vinyl acetate	<u>SU</u>
	1330-20-7	total xylenes	<u>SU</u>

PESTICIDES

CONCENTRATION: LOW MEDIUM HIGH (circle one)
 DATE EXTRACTED/PREPARED: _____
 DATE ANALYZED: _____
 PERCENT MOISTURE: _____
 CONC./DILUTION FACTOR: _____

NOT ANALYZED

PP #	CAS #	NAME	CONC.
(89P)	309-00-2	aldrin	
(90P)	60-57-1	dieldrin	
(91P)	57-74-9	chlordane	
(92P)	50-29-3	4,4'-DDT	
(93P)	72-55-9	4,4'-DDE	
(94P)	72-54-8	4,4'-DDD	
(95P)	115-29-7	α -endosulfan	
(96P)	115-29-7	β -endosulfan	
(97P)	1031-07-8	endosulfan sulfate	
(98P)	72-20-8	endrin	
(99P)	7421-93-4	endrin aldehyde	
(100P)	76-44-8	heptachlor	
(101P)	1024-57-3	heptachlor epoxide	
(102P)	319-84-6	α -BHC	
(103P)	319-85-7	β -BHC	
(104P)	319-86-8	δ -BHC	
(105P)	58-89-9	γ -BHC (lindane)	
(106P)	53469-21-9	PCB-1242	
(107P)	11097-69-1	PCB-1254	
(108P)	11104-28-2	PCB-1221	
(109P)	11141-16-5	PCB-1232	
(110P)	12672-29-6	PCB-1248	
(111P)	11096-82-5	PCB-1260	
(112P)	12674-11-2	PCB-1016	
(113P)	8001-35-2	toxaphene	

DIOXINS

CONCENTRATION: LOW MEDIUM HIGH (circle one)
 DATE EXTRACTED/PREPARED: _____
 DATE ANALYZED: _____
 PERCENT MOISTURE: _____
 CONC./DILUTION FACTOR: _____

PP #	CAS #	NAME	CONC.
(129B)	1746-01-6	2,3,7,8-tetrachlorodibenzo-p-dioxin	

00015

Sample Number
A2753

Laboratory Name: Cambridge Analytical Case No: 2660
 QC Report No: 011

B. Tentatively Identified Compounds

CAS #	Compound Name	Fraction	Scan No. or Retention Time	% Maximum Score Attained Mass Matching Routine: (Specify: _____)	Estimated Concentration (ug/L or ug/kg)
1.	NONE DETECTED	VDA			
2.					
3.					
4.					
5.					
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11.					
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19.					
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21.					
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25.					
26.					
27.					
28.					
29.					
30.					

Sample Number
A2754

Laboratory Name: Cambridge Analytical
QC Report No: 011

Case No: 2660

B. Tentatively Identified Compounds

CAS #	Compound Name	Fraction	Scan No. or Retention Time	% Maximum Score Attained Mass Matching Routine: (Specify: _____)	Estimated Concentration (ug/l or ug/kg)
1.	NONE DETECTED	VOA			
2.					
3.					
4.					
5.					
6.					
7.					
8.					
9.					
10.					
11.					
12.					
13.					
14.					
15.					
16.					
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30.					

A2755

ORGANICS ANALYSIS DATA SHEET

Laboratory Name: Cambridge Analytical Associates
 Lab Sample ID No: 8402-440
 Sample Matrix: Water
 Data Release Authorized By: [Signature]

Case No: 2660
 QC Report No: 011
 Contract No.: 68-01-6791
 Date Sample Received: 5/3/84

VOLATILES

CONCENTRATION: (LOW) MEDIUM HIGH (circle one)
 DATE EXTRACTED/PREPARED: _____
 DATE ANALYZED: 5/5/84
 PERCENT MOISTURE: 100
 CONC./DILUTION FACTOR: 5:1

PP #	CAS #	Compound	ug/l or ug/kg (circle one)
(2V)	107-02-8	acrolein	<u>1000</u>
(3V)	107-13-1	acrylonitrile	<u>1000</u>
(4V)	71-43-2	benzene	<u>SU</u>
(6V)	56-23-5	carbon tetrachloride	<u>SU</u>
(7V)	108-90-7	chlorobenzene	<u>SU</u>
(10V)	107-06-2	1,2-dichloroethane	<u>1U</u>
(11V)	71-55-6	1,1,1-trichloroethane	<u>SU</u>
(13V)	75-34-3	1,1-dichloroethane	<u>SU</u>
(14V)	79-00-5	1,1,2-trichloroethane	<u>SU</u>
(15V)	79-34-5	1,1,2,2-tetrachloroethane	<u>10U</u>
(16V)	75-00-3	chloroethane	<u>10U</u>
(19V)	110-75-8	2-chloroethylvinyl ether	<u>10U</u>
(23V)	67-66-3	chloroform	<u>SU</u>
(29V)	75-35-4	1,1-dichloroethene	<u>SU</u>
(30V)	156-60-5	trans-1,2-dichloroethene	<u>SU</u>
(32V)	78-87-5	1,2-dichloropropane	<u>10U</u>
(33V)	10061-02-6	trans-1,3-dichloropropene	<u>SU</u>
	10061-01-05	cis-1,3-dichloropropene	<u>SU</u>
(38V)	100-41-4	ethylbenzene	<u>SU</u>
(44V)	75-09-2	methylene chloride	<u>SU</u>
(45V)	74-87-3	chloromethane	<u>10U</u>
(46V)	74-83-9	bromomethane	<u>10U</u>
(47V)	75-25-2	bromoform	<u>10U</u>
(48V)	75-27-4	bromodichloromethane	<u>SU</u>
(49V)	75-69-4	fluorotrichloromethane	_____
(50V)	75-71-8	dichlorodifluoromethane	_____
(51V)	124-48-1	chlorodibromomethane	<u>SU</u>
(85V)	127-18-4	tetrachloroethene	<u>SU</u>
(86V)	108-88-3	toluene	<u>SU</u>
(87V)	79-01-6	trichloroethene	<u>SU</u>
(88V)	75-01-4	vinyl chloride	<u>10U</u>
	67-64-1	acetone	<u>SU</u>
	78-93-3	2-butanone	<u>SU</u>
	75-15-0	carbondsulfide	<u>1U</u>
	519-78-6	2-hexanone	<u>SU</u>
	108-10-1	4-methyl-2-pentanone	<u>SU</u>
	100-42-5	styrene	<u>SU</u>
	108-05-4	vinyl acetate	<u>SU</u>
	1330-20-7	total xylenes	<u>SU</u>

PESTICIDES

CONCENTRATION: LOW MEDIUM HIGH (circle one)
 DATE EXTRACTED/PREPARED: _____
 DATE ANALYZED: _____
 PERCENT MOISTURE: _____
 CONC./DILUTION FACTOR: _____

NOT ANALYZED

PP #	CAS #	Compound	ug/l or ug/kg (circle one)
(89P)	309-00-2	aldrin	
(90P)	60-57-1	dieldrin	
(91P)	57-74-9	chlordan	
(92P)	50-29-3	4,4'-DDT	
(93P)	72-55-9	4,4'-DDE	
(94P)	72-54-8	4,4'-DDD	
(95P)	115-29-7	α -endosulfan	
(96P)	115-29-7	β -endosulfan	
(97P)	1031-07-8	endosulfan sulfate	
(98P)	72-20-8	endrin	
(99P)	7421-93-4	endrin aldehyde	
(100P)	76-44-8	heptachlor	
(101P)	1024-57-3	heptachlor epoxide	
(102P)	319-84-6	α -BHC	
(103P)	319-85-7	β -BHC	
(104P)	319-86-8	δ -BHC	
(105P)	58-89-9	γ -BHC (lindane)	
(106P)	53469-21-9	PCB-1242	
(107P)	11097-69-1	PCB-1254	
(108P)	11104-28-2	PCB-1221	
(109P)	11141-16-5	PCB-1232	
(110P)	12672-29-6	PCB-1248	
(111P)	11096-82-5	PCB-1260	
(112P)	12674-11-2	PCB-1016	
(113P)	8001-35-2	toxaphene	

DIOXINS

CONCENTRATION: LOW MEDIUM HIGH (circle one)
 DATE EXTRACTED/PREPARED: _____
 DATE ANALYZED: _____
 PERCENT MOISTURE: _____
 CONC./DILUTION FACTOR: _____

PP #	CAS #	Compound	ug/l or ug/kg (circle one)
(129B)	1746-01-6	2,3,7,8-tetrachlorodibenzo-p-dioxin	

Sample Number
A2755

Laboratory Name: Cambridge Analytical Case No: 2660
QC Report No: 011

B. Tentatively Identified Compounds

CAS #	Compound Name	Fraction	Scan No. or Retention Time	% Maximum Score Attained Mass Matching Routine: (Specify: _____)	Estimated Concentration (ug/L or ug/kg)
1.	NONE DETECTED	VOA			
2.					
3.					
4.					
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2756

ORGANICS ANALYSIS DATA SHEET

Laboratory Name: Cambridge Analytical Associates
Lab Sample ID No: 8402-441
Sample Matrix: Water
Data Release Authorized By: [Signature]

Case No: 2660
QC Report No: 011
Contract No.: 68-01-6791
Date Sample Received: 5/3/84

VOLATILES

CONCENTRATION: LOW MEDIUM HIGH (circle one)
DATE EXTRACTED/PREPARED:
DATE ANALYZED: 5/5/84
PERCENT MOISTURE: 100
CONC./DILUTION FACTOR: 3:1

PESTICIDES

CONCENTRATION: LOW MEDIUM HIGH (circle one)
DATE EXTRACTED/PREPARED:
DATE ANALYZED:
PERCENT MOISTURE:
CONC./DILUTION FACTOR:

Table with columns PP#, CAS#, and chemical name. Includes handwritten values like '100U', 'SU', 'SUR', and 'EL 50 SKR'.

Table with columns PP#, CAS#, and chemical name. Includes handwritten note 'NOT ANALYZED' and values like '100U', 'SU', 'SKR'.

DIOXINS

CONCENTRATION: LOW MEDIUM HIGH (circle one)
DATE EXTRACTED/PREPARED:
DATE ANALYZED:
PERCENT MOISTURE:
CONC./DILUTION FACTOR:

Table with columns PP#, CAS#, and chemical name. Includes handwritten value '00021'.

Sample Number
A2756

Laboratory Name: Cambridge Analytical Case No: 2660
 QC Report No: 011

B. Tentatively Identified Compounds

CAS #	Compound Name	Fraction	Scan No. or Retention Time	% Maximum Score Attained Mass Matching Routine: (Specify: _____)	Estimated Concentration (ug/L or ug/g)
1.	NONE DETECTED	VOA			
2.					
3.					
4.					
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0/22

A2757

ORGANICS ANALYSIS DATA SHEET

Laboratory Name: Cambridge Analytical Associates
Lab Sample ID No: 8402-442
Sample Matrix: Water
Data Release Authorized By: [Signature]

Case No: 2660
QC Report No: 011
Contract No.: 68-01-6791
Date Sample Received: 5/3/84

VOLATILES

CONCENTRATION: (LOW) MEDIUM HIGH (circle one)
DATE EXTRACTED/PREPARED:
DATE ANALYZED: 5/5/84
PERCENT MOISTURE: 100
CONC./DILUTION FACTOR: 3:1

PESTICIDES

CONCENTRATION: LOW MEDIUM HIGH (circle one)
DATE EXTRACTED/PREPARED:
DATE ANALYZED:
PERCENT MOISTURE:
CONC./DILUTION FACTOR:

NOT ANALYZED

Table with columns PP #, CAS #, Name, and Concentration. Includes entries for acrolein, acrylonitrile, benzene, carbon tetrachloride, chlorobenzene, 1,2-dichloroethane, 1,1,1-trichloroethane, 1,1-dichloroethane, 1,1,2-trichloroethane, 1,1,2,2-tetrachloroethane, chloroethane, 2-chloroethylvinyl ether, chloroform, 1,1-dichloroethene, trans-1,2-dichloroethene, 1,2-dichloropropane, trans-1,3-dichloropropene, cis-1,3-dichloropropene, ethylbenzene, methylene chloride, chloromethane, bromomethane, bromoform, bromodichloromethane, fluorotrichloromethane, dichlorodifluoromethane, chlorodibromomethane, tetrachloroethene, toluene, trichloroethene, vinyl chloride, acetone, 2-butanone, carbondisulfide, 2-hexanone, 4-methyl-2-pentanone, styrene, vinyl acetate, total xylenes.

Table with columns PP #, CAS #, Name, and Concentration. Includes entries for aldrin, dieldrin, chlordane, 4,4'-DDT, 4,4'-DDE, 4,4'-DDD, alpha-endosulfan, beta-endosulfan, endosulfan sulfate, endrin, endrin aldehyde, heptachlor, heptachlor epoxide, alpha-BHC, beta-BHC, delta-BHC, gamma-BHC (lindane), PCB-1242, PCB-1254, PCB-1221, PCB-1232, PCB-1248, PCB-1260, PCB-1016, toxaphene.

DIOXINS

CONCENTRATION: LOW MEDIUM HIGH (circle one)
DATE EXTRACTED/PREPARED:
DATE ANALYZED:
PERCENT MOISTURE:
CONC./DILUTION FACTOR:

Table with columns PP #, CAS #, Name, and Concentration. Includes entry for 2,3,7,8-tetrachlorodibenzo-p-dioxin.

00023

Sample Number
A2757

Laboratory Name: Cambridge Analytical
 QC Report No: 011

Case No: 2660

B. Tentatively Identified Compounds

CAS #	Compound Name	Fraction	Scan No. or Retention Time	% Maximum Score Attained Mass Matching Routine: (Specify: _____)	Estimated Concentration (ug/L or ug/kg)
1.	NONE DETECTED	VOA			
2.					
3.					
4.					
5.					
6.					
7.					
8.					
9.					
10.					
11.					
12.					
13.					
14.					
15.					
16.					
17.					
18.					
19.					
20.					
21.					
22.					
23.					
24.					
25.					
26.					
27.					
28.					
29.					
30.					



Cambridge Analytical Associates

Cambridge Analytical Associates, Inc.

KEY TO VOLATILE ORGANICS ANALYSIS

PEAK LABELING

Case Number: 2660

Peak Label	Peak Identity
IS 1	bromochloromethane
IS 2	1-bromo-2-chloropropane
IS 3	1,4-dichlorobutane
SS 1	d ₄ -1,2-dichloroethane
SS 2	d ₈ -toluene
SS 3	4-bromofluorobenzene

00025

RIC
05/05/84 12:04:00
SAMPLE:

DATA: CLPVOA23

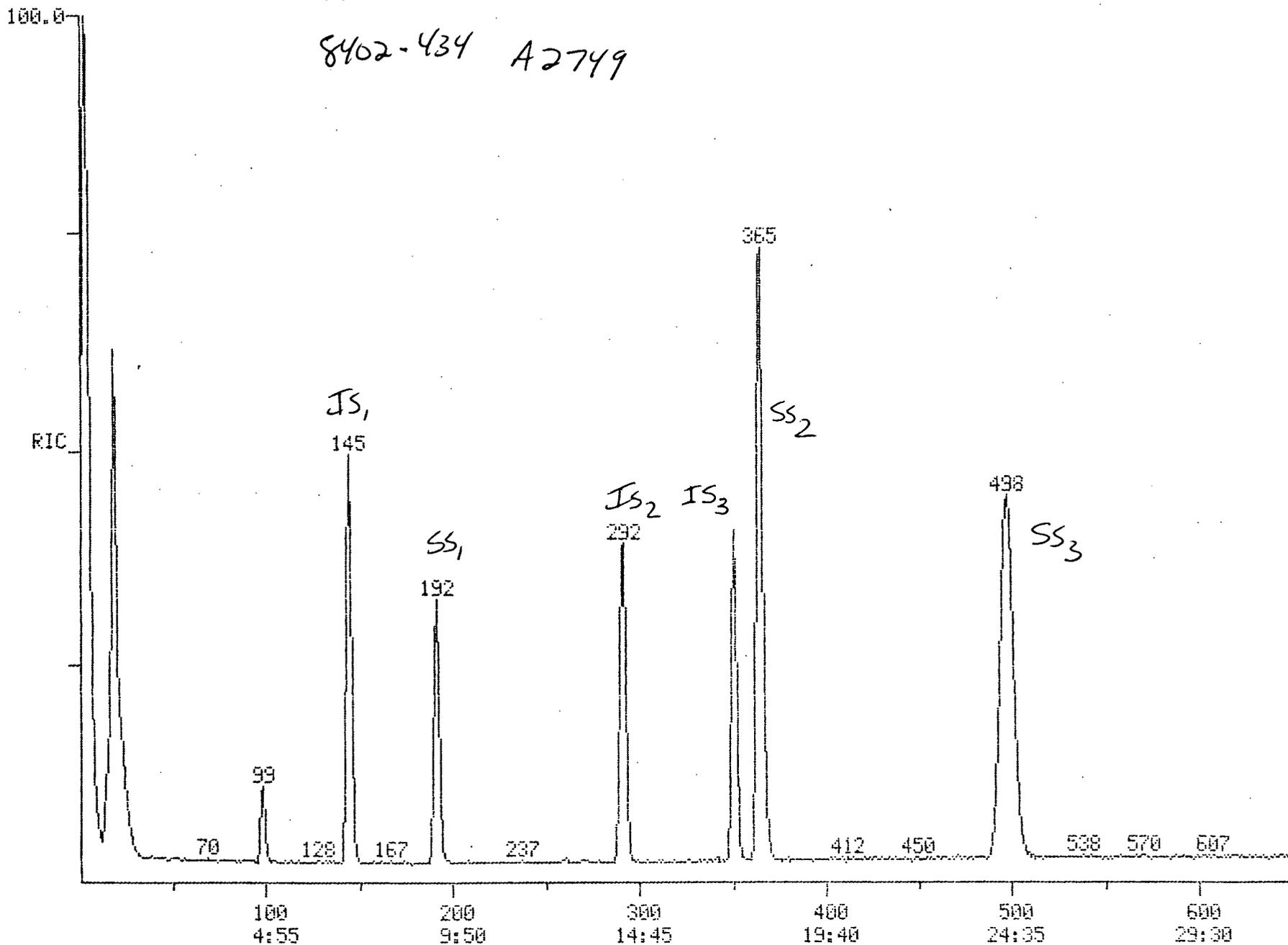
SCANS 1 TO 650

00026

CASE 2660

8402-434 A2749

117376.



A2749 (1984)

SCAN
TIME

FINNIGAN ORGANICS IN WATER ANALYZER
 QUANTITATION REPORT FILE: CLPVOA23

DATA: CLPVOA23.TI
 05/05/84 12:04:00
 SAMPLE:
 SUBMITTED BY:

ANALYST:

AMOUNT=AREA(HGHT) * REF. AMNT/(REF. AREA(HGHT)* RESP. FACT)
 RESP. FAC. FROM LIBRARY ENTRY

- NO NAME
- 1 BROMOCHLOROMETHANE (INTERNAL STANDARD)
- 2 CHLOROMETHANE
- 3 BROMOMETHANE
- 4 VINYL CHLORIDE
- 5 CHLOROETHANE
- 6 METHYLENE CHLORIDE
- 7 CARBON DISULFIDE
- 8 1,1 DICHLOROETHYLENE
- 9 1,1 DICHLOROETHANE
- 10 TRANS 1,2 DICHLOROETHYLENE
- 11 CHLOROFORM
- 12 D4-1,2 DICHLOROETHANE (SURROGATE STANDARD)
- 13 1,2 DICHLOROETHANE
- 14 1,1,1 TRICHLOROETHANE
- 15 CARBON TETRACHLORIDE
- 16 BROMODICHLOROMETHANE
- 17 2-BUTANONE (MEK)
- 18 ACETONE
- 19 ACRYLONITRILE
- 20 ACROLEIN
- 21 1-BROMO-2-CHLOROPROPANE (INTERNAL STANDARD)
- 22 1,2 DICHLOROPROPANE
- 23 TRANS 1,3-DICHLOROPROPENE
- 24 TRICHLOROETHYLENE
- 25 BENZENE
- 26 1,1,2-TRICHLOROETHANE
- 27 CIS 1,3-DICHLOROPROPENE
- 28 DIBROMOCHLOROMETHANE
- 29 1,4 DICHLOROETHANE (INTERNAL STANDARD)
- 30 4-METHYL 2-PENTANONE (MIBK)
- 31 BROMOFORM
- 32 2-HEXANONE (MPK)
- 33 TETRACHLOROETHYLENE
- 34 1,1,2,2 TETRACHLOROETHANE
- 35 D8-TOLUENE (SURROGATE STANDARD)
- 36 TOLUENE
- 37 CHLOROBENZENE
- 38 ETHYLBENZENE
- 39 4-BROMOFLUOROBENZENE (SURROGATE STANDARD)
- 40 STYRENE
- 41 O-XYLENE

NO	M/E	SCAN	TIME	REF	RRT	METH	AREA(HGHT)	AMOUNT	%TOT
1	130	145	7:08	1	1.000	A BB	39589.	50.000 UG/L	11.51
2	NOT FOUND								
3	NOT FOUND								

A2749 (284)

00027

NO	M/E	SCAN	TIME	REF	RRT	METH	AREA(HGHT)	AMOUNT	%TOT
4		NOT FOUND							
5		NOT FOUND							
6	B4	99	4:52	1	0.683	A BB	6918.	2.918 UG/L	0.67 <i>BUK</i>
7		NOT FOUND							
8		NOT FOUND							
9		NOT FOUND							
10		NOT FOUND							
11		NOT FOUND							
12	102	192	9:26	1	1.324	A BB	5412.	82.419 PRCNT	18.98 ✓
13		NOT FOUND							
14		NOT FOUND							
15		NOT FOUND							
16		NOT FOUND							
17		NOT FOUND							
18	43	105	5:10	1	0.724	A BB	552.	3.540 UG/L	0.82 <i>BUK</i>
19		NOT FOUND							
20		NOT FOUND							
21	77	291	14:18	21	1.000	A BB	62414.	50.000 UG/L	11.51
22		NOT FOUND							
23		NOT FOUND							
24		NOT FOUND							
25		NOT FOUND							
26		NOT FOUND							
27		NOT FOUND							
28		NOT FOUND							
29	55	351	17:15	29	1.000	A BB	60354.	50.000 UG/L	11.51
30		NOT FOUND							
31		NOT FOUND							
32		NOT FOUND							
33		NOT FOUND							
34		NOT FOUND							
35	100	364	17:54	29	1.037	A BB	92320.	96.720 PRCNT	22.27 ✓
36	92	368	18:06	29	1.048	A BB	4474.	2.434 UG/L	0.56 <i>BUK</i>
37		NOT FOUND							
38		NOT FOUND							
39	95	498	24:29	29	1.419	A BB	94882.	96.230 PRCNT	22.16 ✓
40		NOT FOUND							
41		NOT FOUND							

A2749 (384)

00028

UNKNOWN SAMPLE QUANTITATION

OWA REVERSE SEARCH STATUS REPORT

THE INTERNAL STANDARD AREA HAS A DIFFERENCE OF: 726 % OF THE LAST STANDARD RUN COMPOUNDS QUANTITATED BY INTERNAL STANDARD USING PARAMETERS IN LIBRARYUX

REPORT ENTRY NO.	EXPECTED SCAN	BEST SCAN	FIT	PURITY	LIBRARY ENTRY NO.	PEAKS FOUND	PEAKS QUANT	SATURATED PEAKS
1	147	145	981	912	1	1	1	0
6	100	99	970	752	6	1	1	0
12	194	192	998	748	12	1	1	0
18	107	105	760	68	18	1	1	0

THE INTERNAL STANDARD AREA HAS A DIFFERENCE OF: 8985 % OF THE LAST STANDARD RUN COMPOUNDS QUANTITATED BY INTERNAL STANDARD USING PARAMETERS IN LIBRARYUY

REPORT ENTRY NO.	EXPECTED SCAN	BEST SCAN	FIT	PURITY	LIBRARY ENTRY NO.	PEAKS FOUND	PEAKS QUANT	SATURATED PEAKS
21	294	291	980	487	1	1	1	0

THE INTERNAL STANDARD AREA HAS A DIFFERENCE OF: -50 % OF THE LAST STANDARD RUN COMPOUNDS QUANTITATED BY INTERNAL STANDARD USING PARAMETERS IN LIBRARYUZ

REPORT ENTRY NO.	EXPECTED SCAN	BEST SCAN	FIT	PURITY	LIBRARY ENTRY NO.	PEAKS FOUND	PEAKS QUANT	SATURATED PEAKS
29	352	351	992	505	1	1	1	0
COMPOUND PASSED REVERSE SEARCH BUT WAS BELOW MINIMUM AREA FOR ION QUANTIT								
32	320	319	504	82	4	1	0	0
35	365	364	971	848	7	1	1	0
36	368	368	902	372	8	1	1	0
39	496	498	987	828	11	1	1	0

NUMBER OF COMPOUNDS IDENTIFIED 10

DATA PROCESSING OF CLPVOA23 COMPLETED ON 5/05/84 13:12:19

A2749(484)

RIC
05/05/84 15:13:00
SAMPLE:

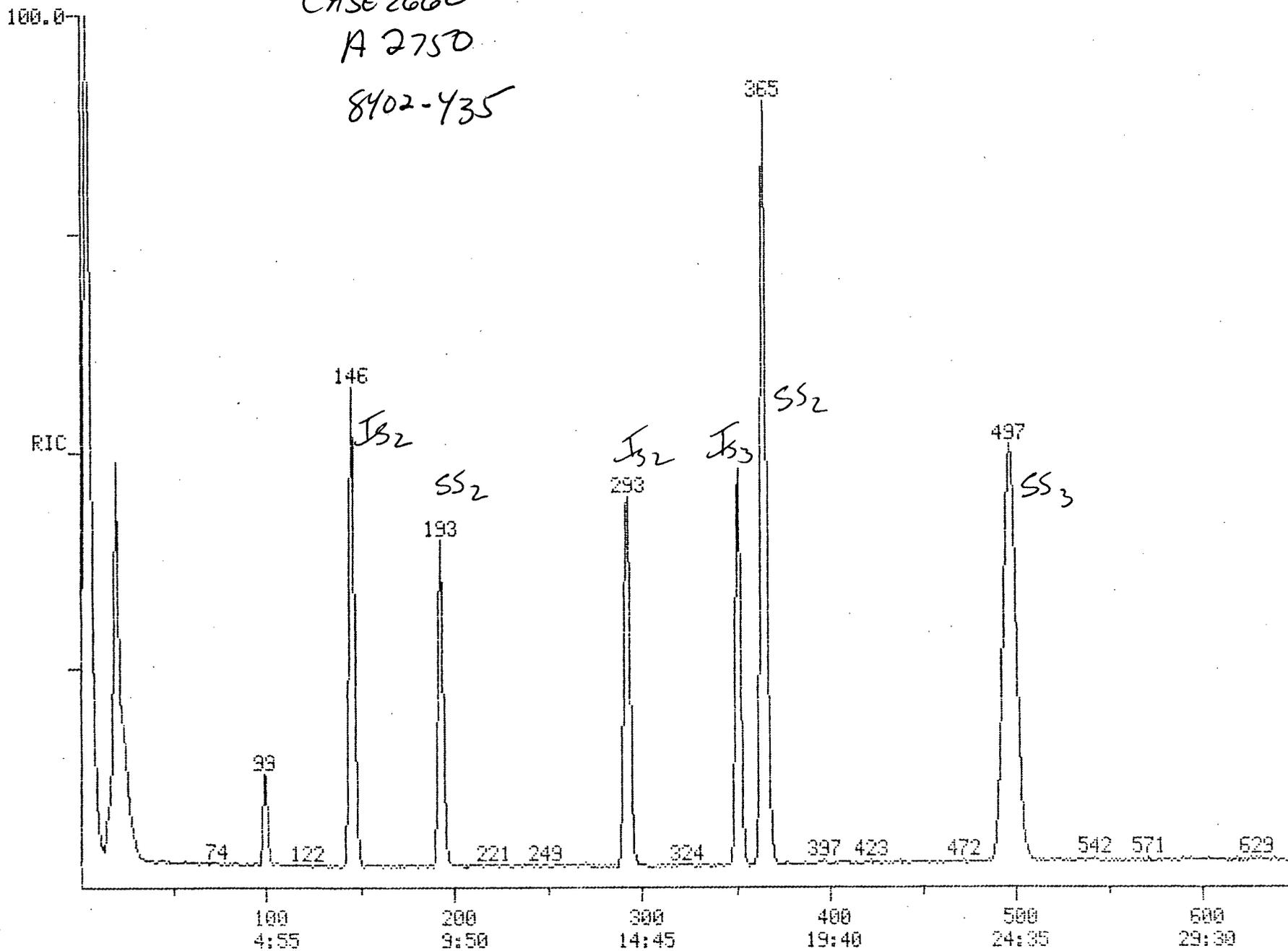
DATA: CLPVOA27

SCANS 1 TO 650

00030

CASE 2660
A 2750
8402-435

93312.



A2750 (1984)

SCAN
TIME

FINNIGAN ORGANICS IN WATER ANALYZER
QUANTITATION REPORT FILE: CLPVOA27

DATA: CLPVOA27.TI
05/05/84 15:13:00
SAMPLE:
SUBMITTED BY:

ANALYST:

AMOUNT=AREA(HGHT) * REF. AMNT / (REF. AREA(HGHT) * RESP. FACT)
RESP. FAC. FROM LIBRARY ENTRY

NO	NAME
1	BROMOCHLOROMETHANE (INTERNAL STANDARD)
2	CHLOROMETHANE
3	BROMOMETHANE
4	VINYL CHLORIDE
5	CHLOROETHANE
6	METHYLENE CHLORIDE
7	CARBON DISULFIDE
8	1,1 DICHLOROETHYLENE
9	1,1 DICHLOROETHANE
10	TRANS 1,2 DICHLOROETHYLENE
11	CHLOROFORM
12	D4-1,2 DICHLOROETHANE (SURROGATE STANDARD)
13	1,2 DICHLOROETHANE
14	1,1,1 TRICHLOROETHANE
15	CARBON TETRACHLORIDE
16	BROMODICHLOROMETHANE
17	2-BUTANONE (MEK)
18	ACETONE
19	ACRYLONITRILE
20	ACROLEIN
21	1-BROMO-2-CHLOROPROPANE (INTERNAL STANDARD)
22	1,2 DICHLOROPROPANE
23	TRANS 1,3-DICHLOROPROPENE
24	TRICHLOROETHYLENE
25	BENZENE
26	1,1,2-TRICHLOROETHANE
27	CIS 1,3-DICHLOROPROPENE
28	DIBROMOCHLOROMETHANE
29	1,4 DICHLOROBUTANE (INTERNAL STANDARD)
30	4-METHYL 2-PENTANONE (MIBK)
31	BROMOFORM
32	2-HEXANONE (MPK)
33	TETRACHLOROETHYLENE
34	1,1,2,2 TETRACHLOROETHANE
35	D8-TOLUENE (SURROGATE STANDARD)
36	TOLUENE
37	CHLOROBENZENE
38	ETHYLBENZENE
39	4-BROMOFLUOROBENZENE (SURROGATE STANDARD)
40	STYRENE
41	O-XYLENE

NO	M/E	SCAN	TIME	REF	RRT	METH	AREA(HGHT)	AMOUNT	%TOT
1	130	146	7:11	1	1.000	A BB	36900.	50.000 UG/L	11.77
2	NOT FOUND								
3	NOT FOUND								

A2750 (284)

00031

NO	M/E	SCAN	TIME	REF	RRT	METH	AREA(HGHT)	AMOUNT	%TOT	
4	NOT	FOUND								
5	NOT	FOUND								
6	84	99	4:52	1	0.678	A BE	6704.	3.034 UG/L	0.71	<i>BK</i>
7	NOT	FOUND								
8	NOT	FOUND								
9	NOT	FOUND								
10	NOT	FOUND								
11	NOT	FOUND								
12	102	193	9:29	1	1.322	A BE	5167.	84.422 PRCNT	19.87	✓
13	NOT	FOUND								
14	NOT	FOUND								
15	NOT	FOUND								
16	NOT	FOUND								
17	NOT	FOUND								
18	43	106	5:13	1	0.726	A BE	583.	4.012 UG/L	0.94	<i>BK</i>
19	NOT	FOUND								
20	NOT	FOUND								
21	77	293	14:24	21	1.000	A BE	57636.	50.000 UG/L	11.77	
22	NOT	FOUND								
23	NOT	FOUND								
24	NOT	FOUND								
25	NOT	FOUND								
26	NOT	FOUND								
27	NOT	FOUND								
28	NOT	FOUND								
29	55	352	17:18	29	1.000	A BE	59016.	50.000 UG/L	11.77	
30	NOT	FOUND								
31	NOT	FOUND								
32	NOT	FOUND								
33	NOT	FOUND								
34	NOT	FOUND								
35	100	365	17:57	29	1.037	A BE	86768.	92.964 PRCNT	21.88	✓
36	NOT	FOUND								
37	NOT	FOUND								
38	NOT	FOUND								
39	95	497	24:26	29	1.412	A BE	87224.	90.470 PRCNT	21.29	✓
40	NOT	FOUND								
41	NOT	FOUND								

A2750 (304)

00032

THE INTERNAL STANDARD AREA HAS A DIFFERENCE OF: 670 % OF THE LAST STANDARD RUN
COMPOUNDS QUANTITATED BY INTERNAL STANDARD USING PARAMETERS IN LIBRARYUX

REPORT ENTRY NO.	EXPECTED SCAN	BEST SCAN	FIT	PURITY	LIBRARY ENTRY NO.	PEAKS FOUND	PEAKS QUANT	SATURATED PEAKS
1	147	146	986	918	1	1	1	0
6	100	99	970	772	6	1	1	0
12	194	193	997	768	12	1	1	0
18	107	106	760	73	18	1	1	0

THE INTERNAL STANDARD AREA HAS A DIFFERENCE OF: 8289 % OF THE LAST STANDARD RUN
COMPOUNDS QUANTITATED BY INTERNAL STANDARD USING PARAMETERS IN LIBRARYUY

REPORT ENTRY NO.	EXPECTED SCAN	BEST SCAN	FIT	PURITY	LIBRARY ENTRY NO.	PEAKS FOUND	PEAKS QUANT	SATURATED PEAKS
21	294	293	977	482	1	1	1	0

THE INTERNAL STANDARD AREA HAS A DIFFERENCE OF: -51 % OF THE LAST STANDARD RUN
COMPOUNDS QUANTITATED BY INTERNAL STANDARD USING PARAMETERS IN LIBRARYUZ

REPORT ENTRY NO.	EXPECTED SCAN	BEST SCAN	FIT	PURITY	LIBRARY ENTRY NO.	PEAKS FOUND	PEAKS QUANT	SATURATED PEAKS
29	352	352	986	508	1	1	1	0
35	365	365	972	848	7	1	1	0
39	496	497	987	836	11	1	1	0

NUMBER OF COMPOUNDS IDENTIFIED B

DATA PROCESSING OF CLPV0A27 COMPLETED ON 5/05/84 16:19:27

A2750 (4084)

00033

RIC
05/05/84 16:00:00
SAMPLE:

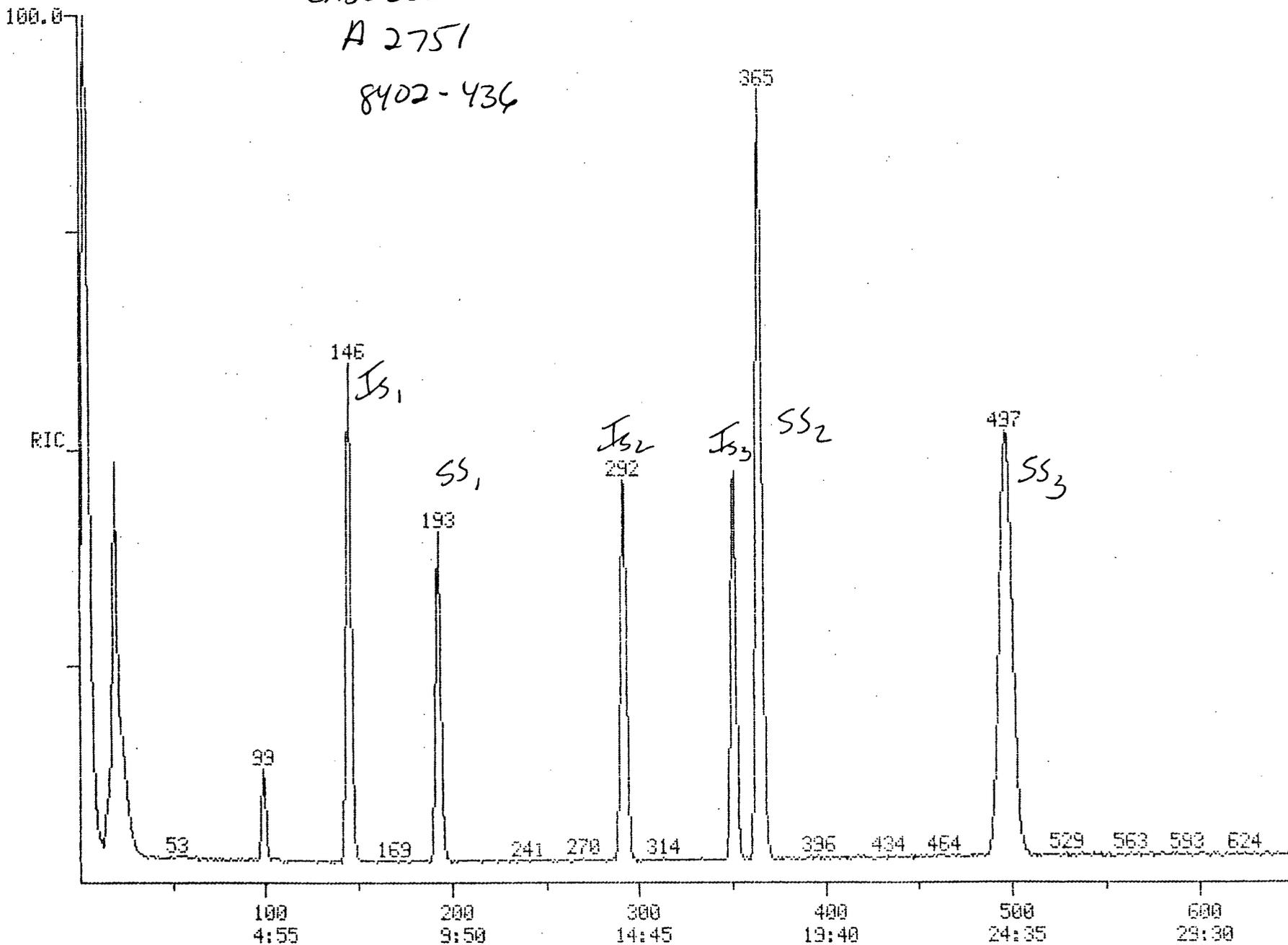
DATA: CLPVOA28

SCANS 1 TO 650

00034

CASE 2660
A 2751
8402-436

91904.



A2751 (1084)

SCAN
TIME

FINNIGAN ORGANICS IN WATER ANALYZER
 QUANTITATION REPORT FILE: CLPVOA28

DATA: CLPVOA28.TI
 05/05/84 16:00:00
 SAMPLE:
 SUBMITTED BY:

ANALYST:

AMOUNT=AREA(HGHT) * REF. AMNT/(REF. AREA(HGHT)* RESP. FACT)
 RESP. FAC. FROM LIBRARY ENTRY

- NO NAME
- 1 BROMOCHLOROMETHANE (INTERNAL STANDARD)
- 2 CHLOROMETHANE
- 3 BROMOMETHANE
- 4 VINYL CHLORIDE
- 5 CHLOROETHANE
- 6 METHYLENE CHLORIDE
- 7 CARBON DISULFIDE
- 8 1,1 DICHLOROETHYLENE
- 9 1,1 DICHLOROETHANE
- 10 TRANS 1,2 DICHLOROETHYLENE
- 11 CHLOROFORM
- 12 D4-1,2 DICHLOROETHANE (SURROGATE STANDARD)
- 13 1,2 DICHLOROETHANE
- 14 1,1,1 TRICHLOROETHANE
- 15 CARBON TETRACHLORIDE
- 16 BROMODICHLOROMETHANE
- 17 2-BUTANONE (MEK)
- 18 ACETONE
- 19 ACRYLONITRILE
- 20 ACROLEIN
- 21 1-BROMO-2-CHLOROPROPANE (INTERNAL STANDARD)
- 22 1,2 DICHLOROPROPANE
- 23 TRANS 1,3-DICHLOROPROPENE
- 24 TRICHLOROETHYLENE
- 25 BENZENE
- 26 1,1,2-TRICHLOROETHANE
- 27 CIS 1,3-DICHLOROPROPENE
- 28 DIBROMOCHLOROMETHANE
- 29 1,4 DICHLOROBUTANE (INTERNAL STANDARD)
- 30 4-METHYL 2-PENTANONE (MIBK)
- 31 BROMOFORM
- 32 2-HEXANONE (MPK)
- 33 TETRACHLOROETHYLENE
- 34 1,1,2,2 TETRACHLOROETHANE
- 35 D8-TOLUENE (SURROGATE STANDARD)
- 36 TOLUENE
- 37 CHLOROBENZENE
- 38 ETHYLBENZENE
- 39 4-BROMOFLUOROBENZENE (SURROGATE STANDARD)
- 40 STYRENE
- 41 O-XYLENE

NO	M/E	SCAN	TIME	REF	RRT	METH	AREA(HGHT)	AMOUNT	%TOT
1	130	146	7:11	1	1.000	A BB	36916.	50.000 UG/L	11.83
2	NOT FOUND								
3	NOT FOUND								

A2757 (2084)

00035

NO	M/E	SCAN	TIME	REF	RRT	METH	AREA(HGHT)	AMOUNT	%TOT
4	NOT FOUND								
5	NOT FOUND								
6	84	99	4:52	1	0.678	A BB	6381.	2.887 UG/L	0.68 <i>BUK</i>
7	NOT FOUND								
8	NOT FOUND								
9	NOT FOUND								
10	NOT FOUND								
11	NOT FOUND								
12	102	193	9:29	1	1.322	A BB	4906.	80.123 PRCNT	18.96 ✓
13	NOT FOUND								
14	NOT FOUND								
15	NOT FOUND								
16	NOT FOUND								
17	NOT FOUND								
18	43	106	5:13	1	0.726	A BB	367.	2.524 UG/L	0.60 <i>BUK</i>
19	NOT FOUND								
20	NOT FOUND								
21	77	292	14:21	21	1.000	A BB	58406.	50.000 UG/L	11.83
22	NOT FOUND								
23	NOT FOUND								
24	NOT FOUND								
25	NOT FOUND								
26	NOT FOUND								
27	NOT FOUND								
28	NOT FOUND								
29	55	352	17:18	29	1.000	A BB	58277.	50.000 UG/L	11.83
30	NOT FOUND								
31	NOT FOUND								
32	NOT FOUND								
33	NOT FOUND								
34	NOT FOUND								
35	100	365	17:57	29	1.037	A BB	86737.	94.110 PRCNT	22.27 ✓
36	NOT FOUND								
37	NOT FOUND								
38	NOT FOUND								
39	95	498	24:29	29	1.415	A BB	88409.	92.861 PRCNT	21.98 ✓
40	NOT FOUND								
41	NOT FOUND								

A2757 (384)

00036

THE INTERNAL STANDARD AREA HAS A DIFFERENCE OF: 671 % OF THE LAST STANDARD RUN
COMPOUNDS QUANTITATED BY INTERNAL STANDARD USING PARAMETERS IN LIBRARYUX

REPORT ENTRY NO.	EXPECTED SCAN	BEST SCAN	FIT	PURITY	LIBRARY ENTRY NO.	PEAKS FOUND	PEAKS QUANT	SATURATED PEAKS
1	147	146	985	915	1	1	1	0
6	100	99	969	766	6	1	1	0
12	194	193	997	752	12	1	1	0
18	107	106	760	37	18	1	1	0

THE INTERNAL STANDARD AREA HAS A DIFFERENCE OF: 8401 % OF THE LAST STANDARD RUN
COMPOUNDS QUANTITATED BY INTERNAL STANDARD USING PARAMETERS IN LIBRARYUY

REPORT ENTRY NO.	EXPECTED SCAN	BEST SCAN	FIT	PURITY	LIBRARY ENTRY NO.	PEAKS FOUND	PEAKS QUANT	SATURATED PEAKS
21	294	292	973	491	1	1	1	0

THE INTERNAL STANDARD AREA HAS A DIFFERENCE OF: -52 % OF THE LAST STANDARD RUN
COMPOUNDS QUANTITATED BY INTERNAL STANDARD USING PARAMETERS IN LIBRARYUZ

REPORT ENTRY NO.	EXPECTED SCAN	BEST SCAN	FIT	PURITY	LIBRARY ENTRY NO.	PEAKS FOUND	PEAKS QUANT	SATURATED PEAKS
29	352	352	993	507	1	1	1	0
35	365	365	970	846	7	1	1	0
39	496	497	990	837	11	1	1	0

NUMBER OF COMPOUNDS IDENTIFIED 8
DATA PROCESSING OF CLPV0A28 COMPLETED ON 5/05/84 17:06:41

A2757 (484)

00037

RIC
05/05/84 16:47:00
SAMPLE:

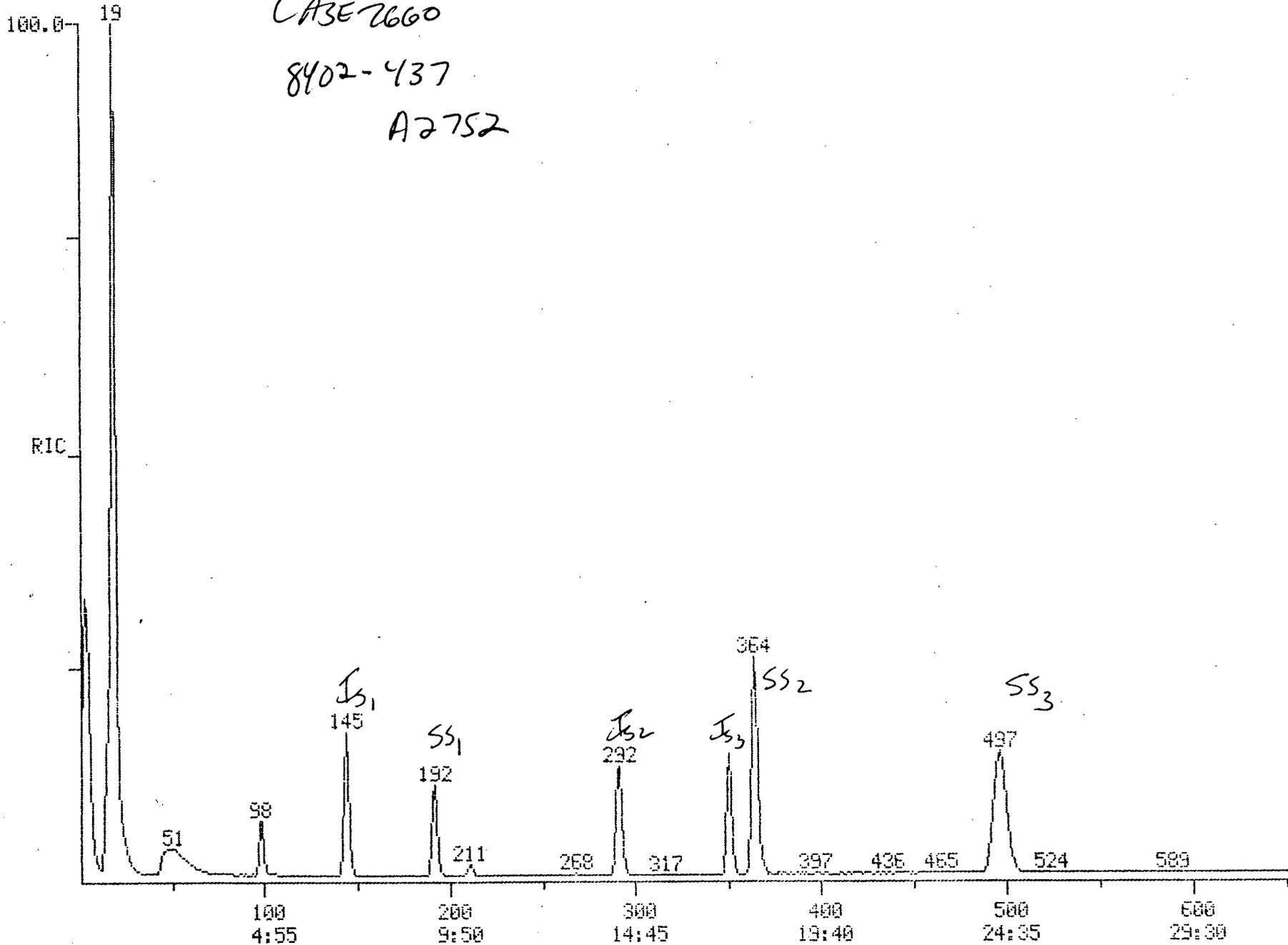
DATA: CLPVOA29

SCANS 1 TO 650

00038

CASE 2660
8402-437
A2752

274432.



A2752 (184)

SCAN
TIME

FINNIGAN ORGANICS IN WATER ANALYZER
QUANTITATION REPORT FILE: CLPVOA29

DATA: CLPVOA29.TI
05/05/84 16:47:00
SAMPLE:
SUBMITTED BY:

ANALYST:

AMOUNT=AREA(HGHT) * REF. AMNT/(REF. AREA(HGHT)* RESP. FACT)
RESP. FAC. FROM LIBRARY ENTRY

- NO NAME
- 1 BROMOCHLOROMETHANE (INTERNAL STANDARD)
- 2 CHLOROMETHANE
- 3 BROMOMETHANE
- 4 VINYL CHLORIDE
- 5 CHLOROETHANE
- 6 METHYLENE CHLORIDE
- 7 CARBON DISULFIDE
- 8 1,1 DICHLOROETHYLENE
- 9 1,1 DICHLOROETHANE
- 10 TRANS 1,2 DICHLOROETHYLENE
- 11 CHLOROFORM
- 12 D4-1,2 DICHLOROETHANE (SURROGATE STANDARD)
- 13 1,2 DICHLOROETHANE
- 14 1,1,1 TRICHLOROETHANE
- 15 CARBON TETRACHLORIDE
- 16 BROMODICHLOROMETHANE
- 17 2-BUTANONE (MEK)
- 18 ACETONE
- 19 ACRYLONITRILE
- 20 ACROLEIN
- 21 1-BROMO-2-CHLOROPROPANE (INTERNAL STANDARD)
- 22 1,2 DICHLOROPROPANE
- 23 TRANS 1,3-DICHLOROPROPENE
- 24 TRICHLOROETHYLENE
- 25 BENZENE
- 26 1,1,2-TRICHLOROETHANE
- 27 CIS 1,3-DICHLOROPROPENE
- 28 DIBROMOCHLOROMETHANE
- 29 1,4 DICHLOROETHANE (INTERNAL STANDARD)
- 30 4-METHYL 2-PENTANONE (MIBK)
- 31 BROMOFORM
- 32 2-HEXANONE (MPK)
- 33 TETRACHLOROETHYLENE
- 34 1,1,2,2 TETRACHLOROETHANE
- 35 DB-TOLUENE (SURROGATE STANDARD)
- 36 TOLUENE
- 37 CHLOROBENZENE
- 38 ETHYLBENZENE
- 39 4-BROMOFLUOROBENZENE (SURROGATE STANDARD)
- 40 STYRENE
- 41 O-XYLENE

NO	M/E	SCAN	TIME	REF	RRT	METH	AREA(HGHT)	AMOUNT	%TOT
1	130	145	7:08	1	1.000	A BB	32366.	50.000 UG/L	11.51
2	NOT FOUND								
3	NOT FOUND								

A2752(284)

00039

NO	M/E	SCAN	TIME	REF	RRT	METH	AREA (HGHT)	AMOUNT	%TOT
4	NOT FOUND								
5	NOT FOUND								
6	84	98	4:49	1	0.676	A BB	11813.	6.095 UG/L	1.40 <i>BLK</i>
7	NOT FOUND								
8	NOT FOUND								
9	NOT FOUND								
10	NOT FOUND								
11	NOT FOUND								
12	102	192	9:26	1	1.324	A BB	4242.	79.018 PRCNT	18.19 ✓
13	NOT FOUND								
14	NOT FOUND								
15	NOT FOUND								
16	NOT FOUND								
17	NOT FOUND								
18	43	105	5:10	1	0.724	A BB	1688.	13.242 UG/L	3.05 <i>BLK</i>
19	NOT FOUND								
20	NOT FOUND								
21	77	292	14:21	21	1.000	A BB	49904.	50.000 UG/L	11.51
22	NOT FOUND								
23	NOT FOUND								
24	NOT FOUND								
25	NOT FOUND								
26	NOT FOUND								
27	NOT FOUND								
28	NOT FOUND								
29	55	351	17:15	29	1.000	A BB	51853.	50.000 UG/L	11.51
30	NOT FOUND								
31	NOT FOUND								
32	NOT FOUND								
33	NOT FOUND								
34	NOT FOUND								
35	100	364	17:54	29	1.037	A BB	75771.	92.396 PRCNT	21.27 ✓
36	92	367	18:03	29	1.046	A BB	5360.	3.394 UG/L	0.78 <i>BLK</i>
37	NOT FOUND								
38	NOT FOUND								
39	95	497	24:26	29	1.416	A BB	76377.	90.162 PRCNT	20.76 ✓
40	NOT FOUND								
41	NOT FOUND								

A2752 (384)

00040

THE INTERNAL STANDARD AREA HAS A DIFFERENCE OF: 575 % OF THE LAST STANDARD RUN
COMPOUNDS QUANTITATED BY INTERNAL STANDARD USING PARAMETERS IN LIBRARYUX

REPORT ENTRY NO.	EXPECTED SCAN	BEST SCAN	FIT	PURITY	LIBRARY ENTRY NO.	PEAKS FOUND	PEAKS QUANT	SATURATED PEAKS
1	147	145	984	911	1	1	1	0
6	100	98	991	865	6	1	1	0
12	194	192	997	756	12	1	1	0
18	107	105	761	159	18	1	1	0

THE INTERNAL STANDARD AREA HAS A DIFFERENCE OF: 7164 % OF THE LAST STANDARD RUN
COMPOUNDS QUANTITATED BY INTERNAL STANDARD USING PARAMETERS IN LIBRARYUY

REPORT ENTRY NO.	EXPECTED SCAN	BEST SCAN	FIT	PURITY	LIBRARY ENTRY NO.	PEAKS FOUND	PEAKS QUANT	SATURATED PEAKS
21	294	292	984	472	1	1	1	0

THE INTERNAL STANDARD AREA HAS A DIFFERENCE OF: -57 % OF THE LAST STANDARD RUN
COMPOUNDS QUANTITATED BY INTERNAL STANDARD USING PARAMETERS IN LIBRARYUZ

REPORT ENTRY NO.	EXPECTED SCAN	BEST SCAN	FIT	PURITY	LIBRARY ENTRY NO.	PEAKS FOUND	PEAKS QUANT	SATURATED PEAKS
29	352	351	991	508	1	1	1	0
35	365	364	969	852	7	1	1	0
36	368	367	852	264	8	1	1	0
39	496	497	989	843	11	1	1	0

NUMBER OF COMPOUNDS IDENTIFIED 9

DATA PROCESSING OF CLPVOA29 COMPLETED ON 5/05/84 17:53:50

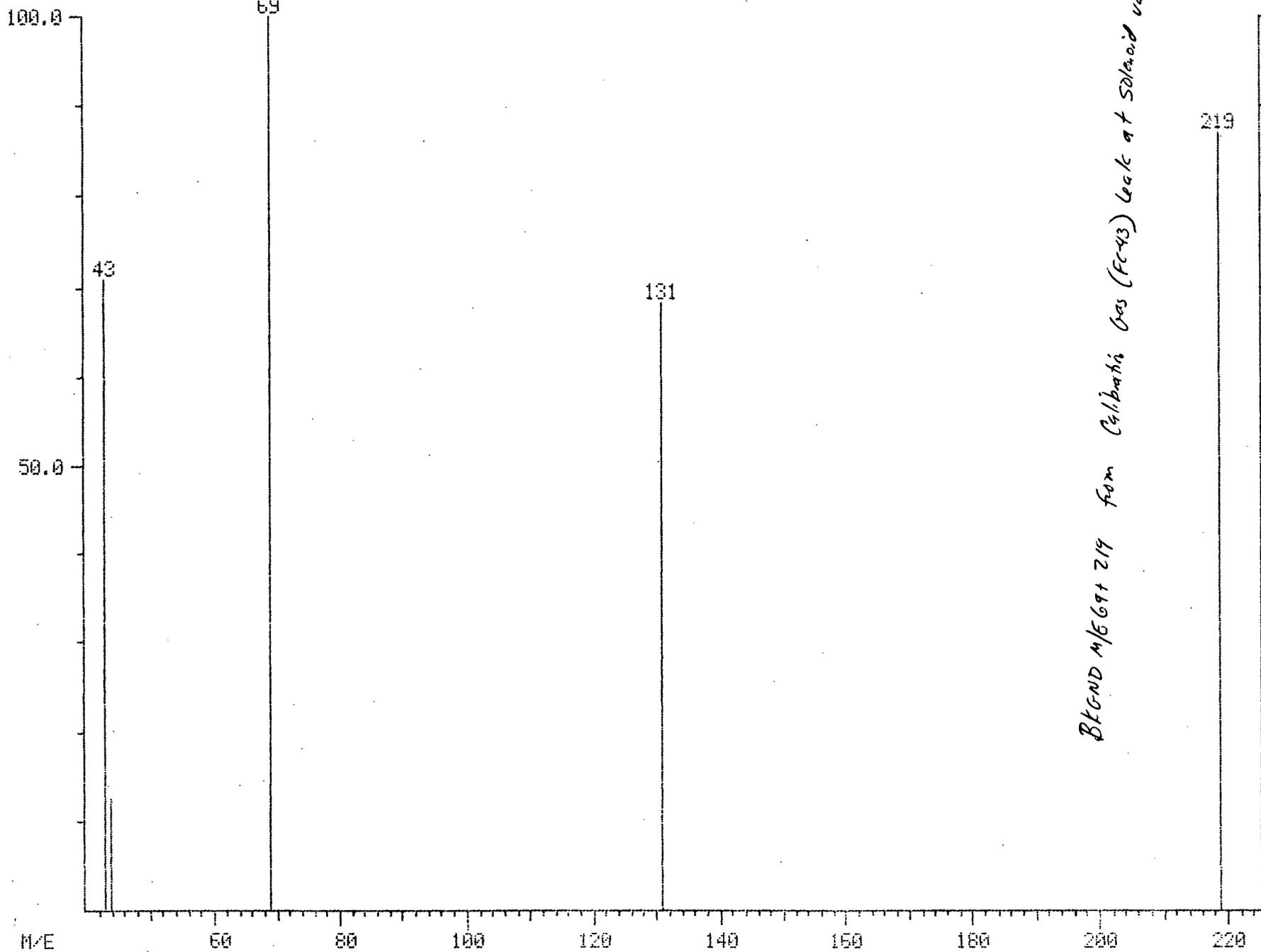
A2752 (4084)

00041

MASS SPECTRUM
05/05/84 16:47:00 + 5:10
SAMPLE:

DATA: CLP00A29 #105

BASE M/E: 69
RIC: 3100.



Blend m/e 69 + 219 from Calibration Gas (FC-43) leak at solvent valve.

A7752 - Acetone

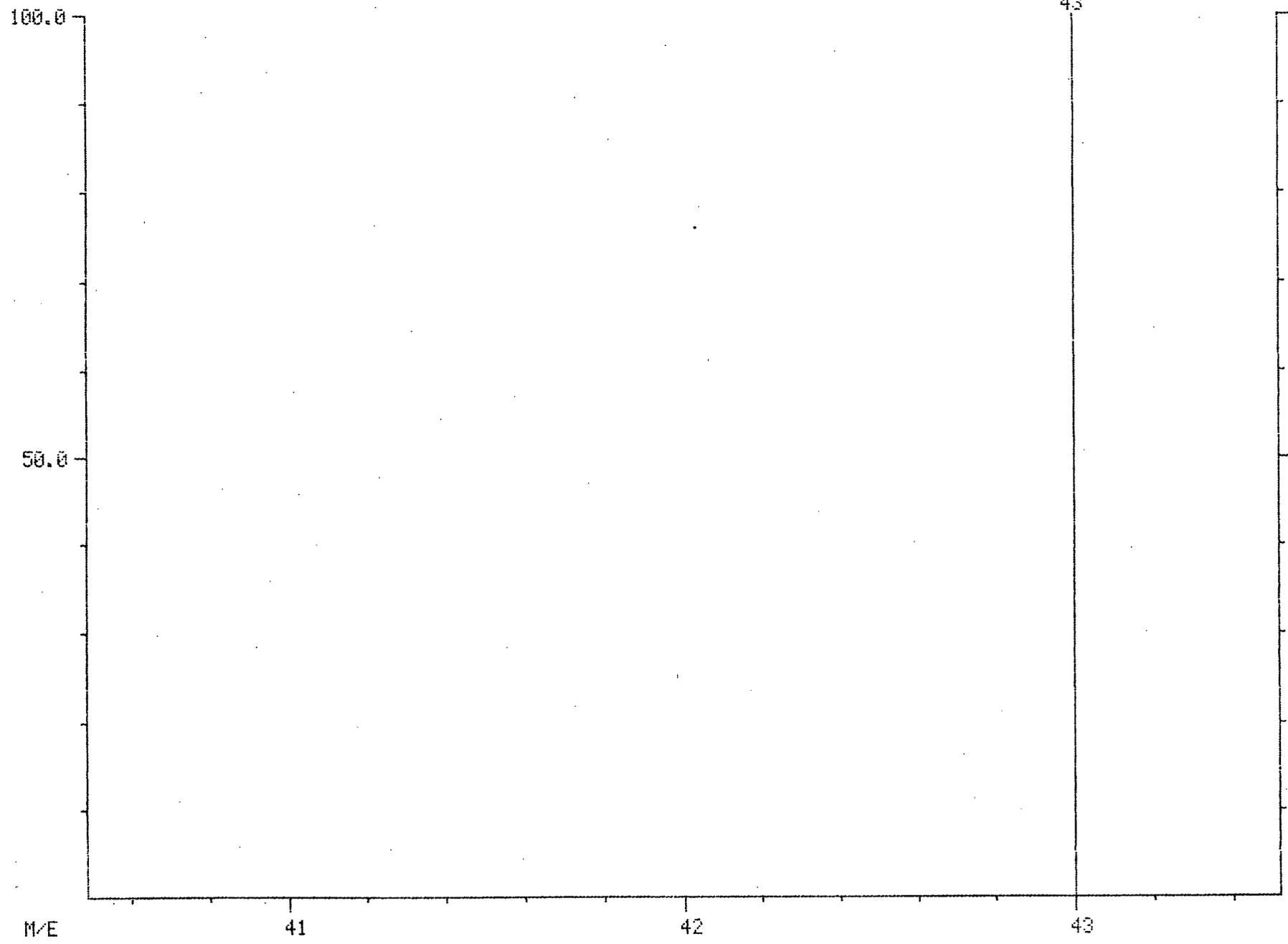
00042

MASS SPECTRUM
05/05/84 16:47:00 + 5:10
SAMPLE:
ENHANCED (S 15B 2N)

DATA: CLP00A29 #105

BASE M/E: 43
RIC: 556.

00043



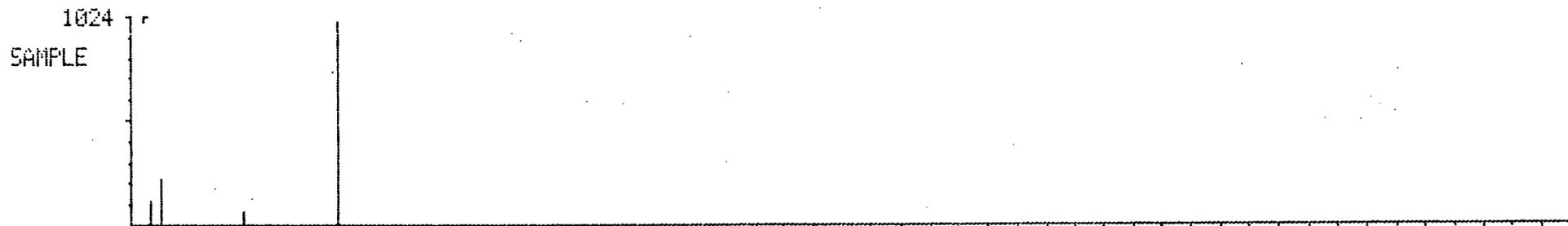
A275a - Acetone

LIBRARY SEARCH
05/05/84 16:47:00 + 10:22
SAMPLE:
ENHANCED (S 15B 2N 0T)

DATA: CLP00A29 # 211

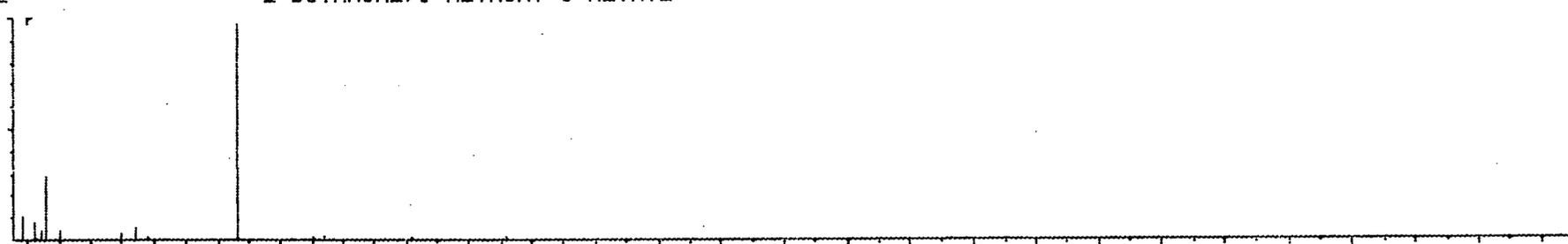
BASE M/E: 73
RIC: 3357.

00044
A2752 - Tentative ID Cpd.



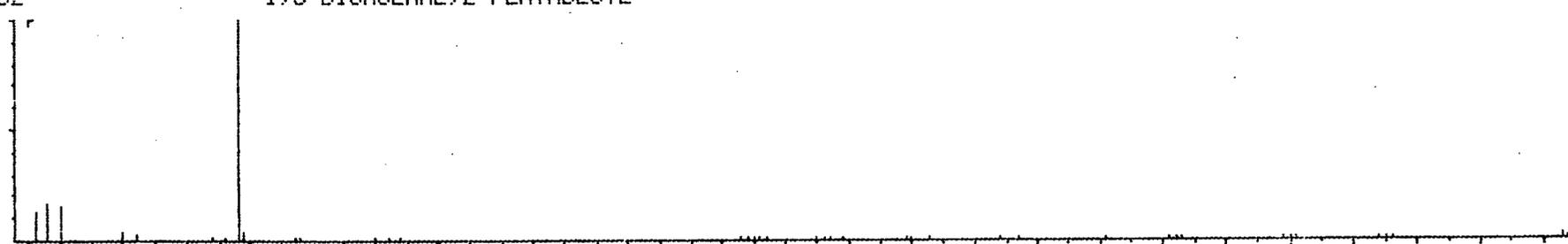
06.H12.02
M WT 116
B PK 73
RANK 1
IN 2220
PUR 914

2-BUTANONE, 3-METHOXY-3-METHYL-



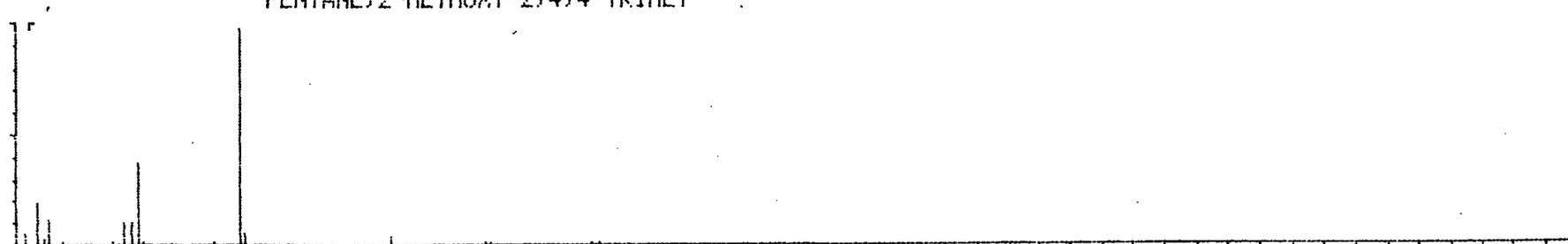
018.H36.02
M WT 284
B PK 73
RANK 2
IN 20870
PUR 747

1,3-DIOXOLANE, 2-PENTADECYL-



09.H20.0
M WT 144
B PK 73
RANK 3
IN 5296
PUR 739

PENTANE, 2-METHOXY-2,4,4-TRIMET



M/E 50 100 150 200 250

RIC
05/05/84 17:34:00
SAMPLE:

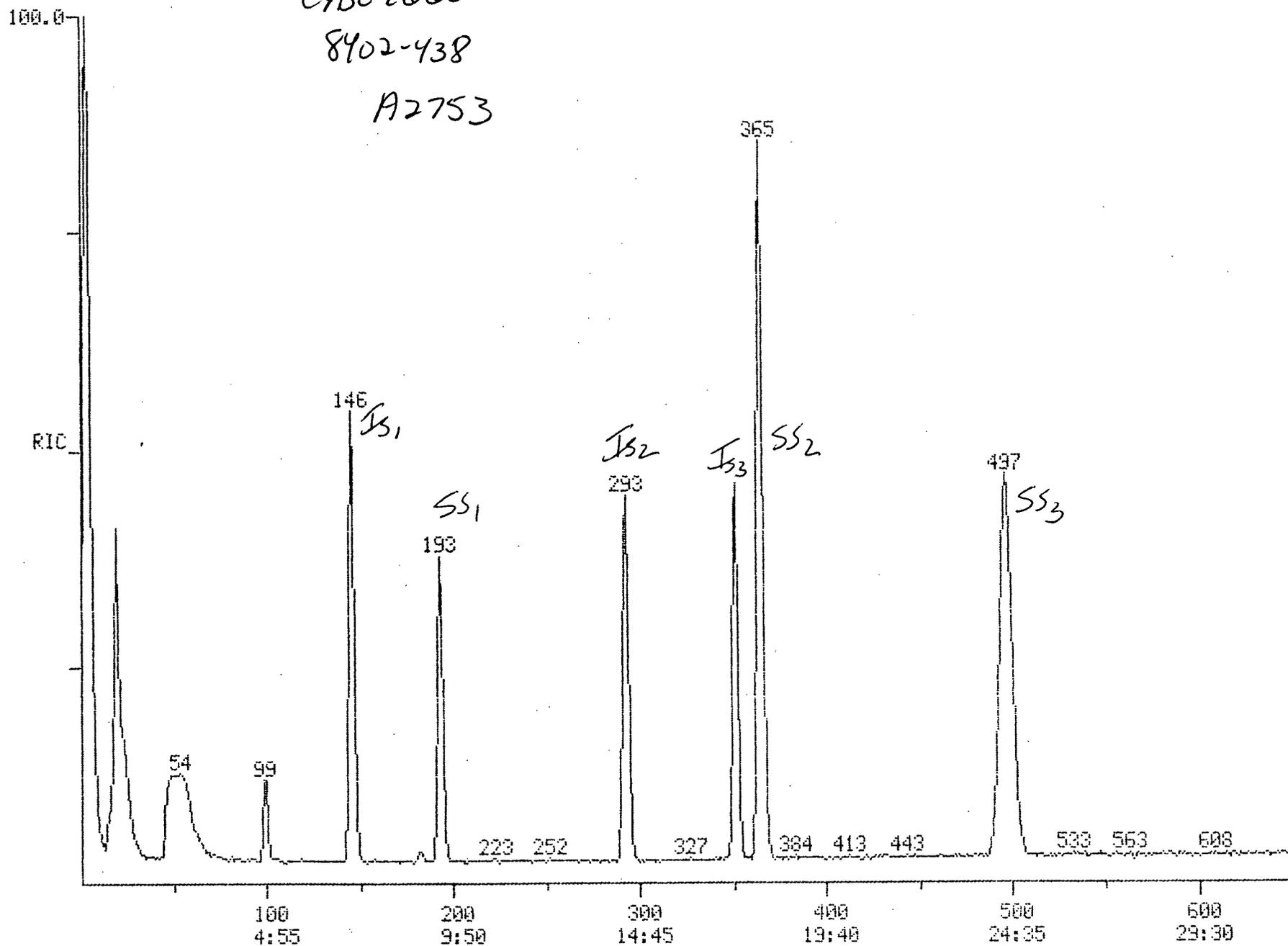
DATA: CLPVOA30

SCANS 1 TO 650

00045

CASE 2060
8402-438
A2753

87424.



A2753 (1084)

SCAN
TIME

FINNIGAN ORGANICS IN WATER ANALYZER
 QUANTITATION REPORT FILE: CLPVOA30

DATA: CLPVOA30.TI
 05/05/84 17:34:00
 SAMPLE:
 SUBMITTED BY:

ANALYST:

AMOUNT=AREA(HGHT) * REF. AMNT / (REF. AREA(HGHT) * RESP. FACT)
 RESP. FAC. FROM LIBRARY ENTRY

- NO NAME
- 1 BROMOCHLOROMETHANE (INTERNAL STANDARD)
- 2 CHLOROMETHANE
- 3 BROMOMETHANE
- 4 VINYL CHLORIDE
- 5 CHLOROETHANE
- 6 METHYLENE CHLORIDE
- 7 CARBON DISULFIDE
- 8 1, 1 DICHLOROETHYLENE
- 9 1, 1 DICHLOROETHANE
- 10 TRANS 1, 2 DICHLOROETHYLENE
- 11 CHLOROFORM
- 12 D4-1, 2 DICHLOROETHANE (SURROGATE STANDARD)
- 13 1, 2 DICHLOROETHANE
- 14 1, 1, 1 TRICHLOROETHANE
- 15 CARBON TETRACHLORIDE
- 16 BROMODICHLOROMETHANE
- 17 2-BUTANONE (MEK)
- 18 ACETONE
- 19 ACRYLONITRILE
- 20 ACROLEIN
- 21 1-BROMO-2-CHLOROPROPANE (INTERNAL STANDARD)
- 22 1, 2 DICHLOROPROPANE
- 23 TRANS 1, 3-DICHLOROPROPENE
- 24 TRICHLOROETHYLENE
- 25 BENZENE
- 26 1, 1, 2-TRICHLOROETHANE
- 27 CIS 1, 3-DICHLOROPROPENE
- 28 DIBROMOCHLOROMETHANE
- 29 1, 4 DICHLOROBUTANE (INTERNAL STANDARD)
- 30 4-METHYL 2-PENTANONE (MIBK)
- 31 BROMOFORM
- 32 2-HEXANONE (MPK)
- 33 TETRACHLOROETHYLENE
- 34 1, 1, 2, 2 TETRACHLOROETHANE
- 35 D8-TOLUENE (SURROGATE STANDARD)
- 36 TOLUENE
- 37 CHLOROBENZENE
- 38 ETHYLBENZENE
- 39 4-BROMOFLUOROBENZENE (SURROGATE STANDARD)
- 40 STYRENE
- 41 O-XYLENE

NO	M/E	SCAN	TIME	REF	RRT	METH	AREA(HGHT)	AMOUNT	%TOT
1	130	146	7:11	1	1.000	A BB	33022.	50.000 UG/L	12.07
2	NOT FOUND								
3	NOT FOUND								

A2753 (284)

00046

NO	M/E	SCAN	TIME	REF	RRT	METH	AREA(HGHT)	AMOUNT	%TOT
4		NOT FOUND							
5		NOT FOUND							
6	84	100	4:55	1	0.685	A BB	6072.	3.071 UG/L	0.74 <i>BLK</i>
7		NOT FOUND							
8		NOT FOUND							
9		NOT FOUND							
10		NOT FOUND							
11		NOT FOUND							
12	102	193	9:29	1	1.322	A BB	4457.	81.374 PRONT	19.65 ✓
13		NOT FOUND							
14		NOT FOUND							
15		NOT FOUND							
16		NOT FOUND							
17		NOT FOUND							
18	43	106	5:13	1	0.726	A BB	305.	2.345 UG/L	0.57 <i>BLK</i>
19		NOT FOUND							
20		NOT FOUND							
21	77	293	14:24	21	1.000	A BB	52572.	50.000 UG/L	12.07
22		NOT FOUND							
23		NOT FOUND							
24		NOT FOUND							
25		NOT FOUND							
26		NOT FOUND							
27		NOT FOUND							
28		NOT FOUND							
29	55	352	17:18	29	1.000	A BB	53385.	50.000 UG/L	12.07
30		NOT FOUND							
31		NOT FOUND							
32		NOT FOUND							
33		NOT FOUND							
34		NOT FOUND							
35	100	365	17:57	29	1.037	A BB	77145.	91.372 PRONT	22.06 ✓
36		NOT FOUND							
37		NOT FOUND							
38		NOT FOUND							
39	95	497	24:26	29	1.412	A BB	75012.	86.010 PRONT	20.77 ✓
40		NOT FOUND							
41		NOT FOUND							

A2753 (384)

00047

THE INTERNAL STANDARD AREA HAS A DIFFERENCE OF: 589 % OF THE LAST STANDARD RUN
COMPOUNDS QUANTITATED BY INTERNAL STANDARD USING PARAMETERS IN LIBRARYUX

REPORT ENTRY NO.	EXPECTED SCAN	BEST SCAN	FIT	PURITY	LIBRARY ENTRY NO.	PEAKS FOUND	PEAKS QUANT	SATURATED PEAKS
1	147	146	980	909	1	1	1	0
6	100	100	964	772	6	1	1	0
12	194	193	1000	765	12	1	1	0
18	107	106	759	50	18	1	1	0

THE INTERNAL STANDARD AREA HAS A DIFFERENCE OF: 7552 % OF THE LAST STANDARD RUN
COMPOUNDS QUANTITATED BY INTERNAL STANDARD USING PARAMETERS IN LIBRARYUY

REPORT ENTRY NO.	EXPECTED SCAN	BEST SCAN	FIT	PURITY	LIBRARY ENTRY NO.	PEAKS FOUND	PEAKS QUANT	SATURATED PEAKS
21	294	293	977	484	1	1	1	0

THE INTERNAL STANDARD AREA HAS A DIFFERENCE OF: -56 % OF THE LAST STANDARD RUN
COMPOUNDS QUANTITATED BY INTERNAL STANDARD USING PARAMETERS IN LIBRARYUZ

REPORT ENTRY NO.	EXPECTED SCAN	BEST SCAN	FIT	PURITY	LIBRARY ENTRY NO.	PEAKS FOUND	PEAKS QUANT	SATURATED PEAKS
29	352	352	988	513	1	1	1	0
35	365	365	970	850	7	1	1	0
39	496	497	989	833	11	1	1	0

NUMBER OF COMPOUNDS IDENTIFIED 8

DATA PROCESSING OF CLPV0A30 COMPLETED ON 5/05/84 18:40:22

A2753 (4084)

00048

RIC
05/05/84 23:59:00
SAMPLE:

DATA: CLPVOA3E

SCANS 1 TO 650

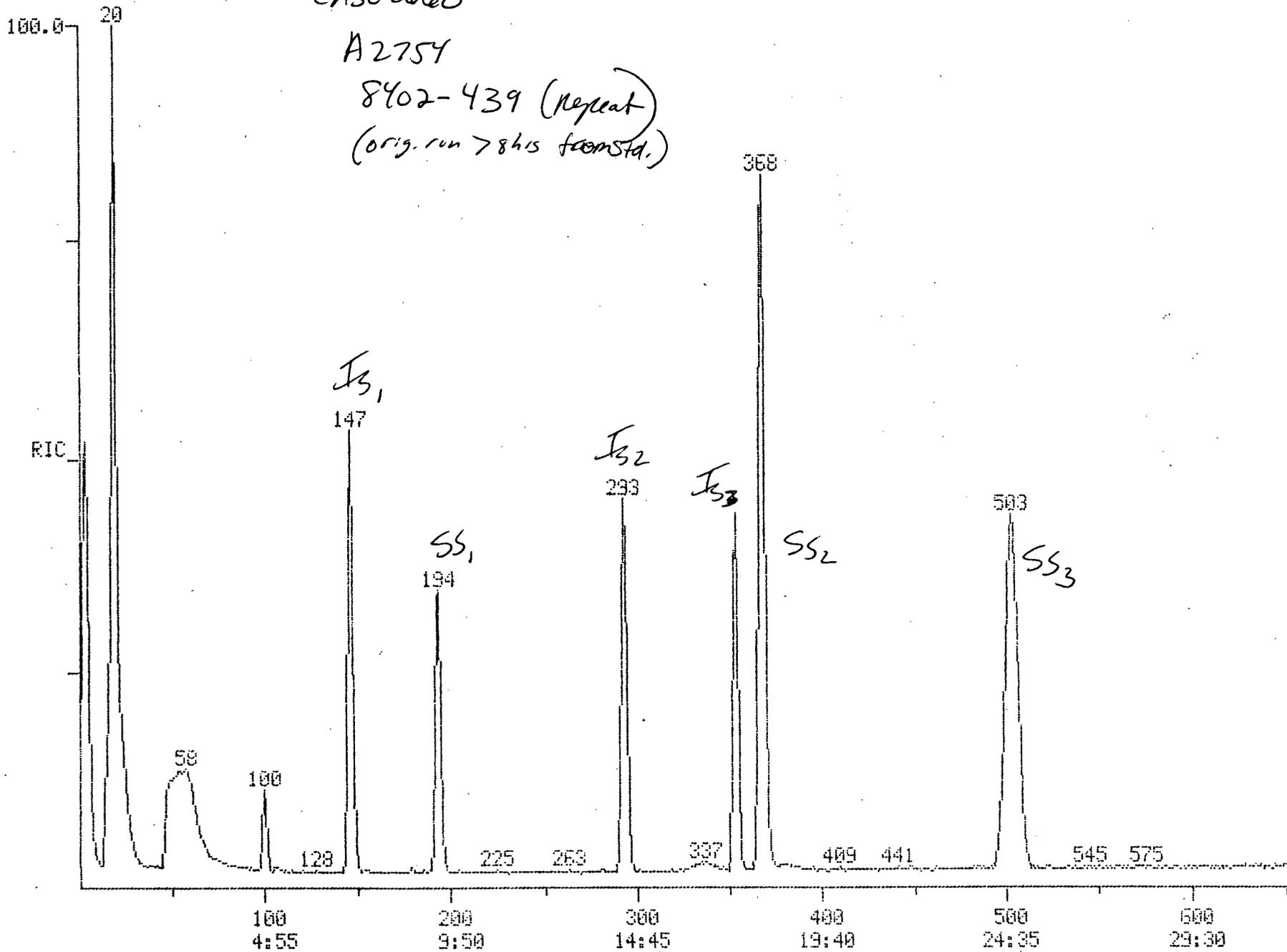
00049

CASE 2660

A2754

8402-439 (Repeat)
(orig. run 78415 from Std.)

83840.



A2754 (1084)

SCAN
TIME

FINNIGAN ORGANICS IN WATER ANALYZER
 QUANTITATION REPORT FILE: CLPVOA36

DATA: CLPVOA36.TI
 05/05/84 23:59:00
 SAMPLE:
 SUBMITTED BY:

ANALYST:

AMOUNT=AREA(HGHT) * REF. AMNT/(REF. AREA(HGHT)* RESP. FACT)
 RESP. FAC. FROM LIBRARY ENTRY

- NO NAME
- 1 BROMOCHLOROMETHANE (INTERNAL STANDARD)
- 2 CHLOROMETHANE
- 3 BROMOMETHANE
- 4 VINYL CHLORIDE
- 5 CHLOROETHANE
- 6 METHYLENE CHLORIDE
- 7 CARBON DISULFIDE
- 8 1,1 DICHLOROETHYLENE
- 9 1,1 DICHLOROETHANE
- 10 TRANS 1,2 DICHLOROETHYLENE
- 11 CHLOROFORM
- 12 D4-1,2 DICHLOROETHANE (SURROGATE STANDARD)
- 13 1,2 DICHLOROETHANE
- 14 1,1,1 TRICHLOROETHANE
- 15 CARBON TETRACHLORIDE
- 16 BROMODICHLOROMETHANE
- 17 2-BUTANONE (MEK)
- 18 ACETONE
- 19 ACRYLONITRILE
- 20 ACROLEIN
- 21 1-BROMO-2-CHLOROPROPANE (INTERNAL STANDARD)
- 22 1,2 DICHLOROPROPANE
- 23 TRANS 1,3-DICHLOROPROPENE
- 24 TRICHLOROETHYLENE
- 25 BENZENE
- 26 1,1,2-TRICHLOROETHANE
- 27 CIS 1,3-DICHLOROPROPENE
- 28 DIBROMOCHLOROMETHANE
- 29 1,4 DICHLOROBUTANE (INTERNAL STANDARD)
- 30 4-METHYL 2-PENTANONE (MIEK)
- 31 BROMOFORM
- 32 2-HEXANONE (MPK)
- 33 TETRACHLOROETHYLENE
- 34 1,1,2,2 TETRACHLOROETHANE
- 35 D8-TOLUENE (SURROGATE STANDARD)
- 36 TOLUENE
- 37 CHLOROBENZENE
- 38 ETHYLBENZENE
- 39 4-BROMOFLUOROBENZENE (SURROGATE STANDARD)
- 40 STYRENE
- 41 O-XYLENE

NO	M/E	SCAN	TIME	REF	RRT	METH	AREA(HGHT)	AMOUNT	%TOT
1	130	147	7:14	1	1.000	A BB	30641.	50.000 UG/L	12.38
2	NOT FOUND								
3	NOT FOUND								

A2754 (2084)

00050

NO	M/E	SCAN	TIME	REF	RRT	METH	AREA(HGHT)	AMOUNT	%TOT
4	NOT	FOUND							
5	NOT	FOUND							
6	84	100	4:55	1	0.680	A BB	5344.	2.363 UG/L	0.59 <i>BUK</i>
7	NOT	FOUND							
8	NOT	FOUND							
9	NOT	FOUND							
10	NOT	FOUND							
11	NOT	FOUND							
12	102	193	9:29	1	1.313	A BB	3845.	84.751 PRCNT	20.98 ✓
13	NOT	FOUND							
14	NOT	FOUND							
15	NOT	FOUND							
16	NOT	FOUND							
17	NOT	FOUND							
18	43	107	5:16	1	0.728	A BB	684.	5.475 UG/L	1.36 <i>BUK</i>
19	NOT	FOUND							
20	NOT	FOUND							
21	77	293	14:24	21	1.000	A BB	50534.	50.000 UG/L	12.38
22	NOT	FOUND							
23	NOT	FOUND							
24	NOT	FOUND							
25	NOT	FOUND							
26	NOT	FOUND							
27	NOT	FOUND							
28	NOT	FOUND							
29	55	354	17:24	29	1.000	A BB	49832.	50.000 UG/L	12.38
30	NOT	FOUND							
31	NOT	FOUND							
32	NOT	FOUND							
33	NOT	FOUND							
34	NOT	FOUND							
35	100	368	18:06	29	1.040	A BB	74861.	77.755 PRCNT	19.25 ✓
36	NOT	FOUND							
37	NOT	FOUND							
38	NOT	FOUND							
39	95	503	24:44	29	1.421	A BB	68413.	83.569 PRCNT	20.69 ✓
40	NOT	FOUND							
41	NOT	FOUND							

A2754 (384)

00051

THE INTERNAL STANDARD AREA HAS A DIFFERENCE OF: 539 % OF THE LAST STANDARD RUN
COMPOUNDS QUANTITATED BY INTERNAL STANDARD USING PARAMETERS IN LIBRARYUX

REPORT ENTRY NO.	EXPECTED SCAN	BEST SCAN	FIT	PURITY	LIBRARY ENTRY NO.	PEAKS FOUND	PEAKS QUANT	SATURATED PEAKS
1	148	147	986	932	1	1	1	0
6	101	100	961	796	6	1	1	0
12	195	194	996	773	12	1	1	0
18	108	107	760	111	18	1	1	0

THE INTERNAL STANDARD AREA HAS A DIFFERENCE OF: 7255 % OF THE LAST STANDARD RUN
COMPOUNDS QUANTITATED BY INTERNAL STANDARD USING PARAMETERS IN LIBRARYUY

REPORT ENTRY NO.	EXPECTED SCAN	BEST SCAN	FIT	PURITY	LIBRARY ENTRY NO.	PEAKS FOUND	PEAKS QUANT	SATURATED PEAKS
21	295	293	974	494	1	1	1	0

THE INTERNAL STANDARD AREA HAS A DIFFERENCE OF: -59 % OF THE LAST STANDARD RUN
COMPOUNDS QUANTITATED BY INTERNAL STANDARD USING PARAMETERS IN LIBRARYUZ

REPORT ENTRY NO.	EXPECTED SCAN	BEST SCAN	FIT	PURITY	LIBRARY ENTRY NO.	PEAKS FOUND	PEAKS QUANT	SATURATED PEAKS
29	355	354	984	528	1	1	1	0
35	368	368	970	855	7	1	1	0
39	506	503	989	863	11	1	1	0

NUMBER OF COMPOUNDS IDENTIFIED 8
DATA PROCESSING OF CLPV0A36 COMPLETED ON 5/06/84 0:41:18

A2754 (484)

00052

RIC
05/05/84 21:49:00
SAMPLE:

DATA: CLPVD033

SCANS 1 TO 650

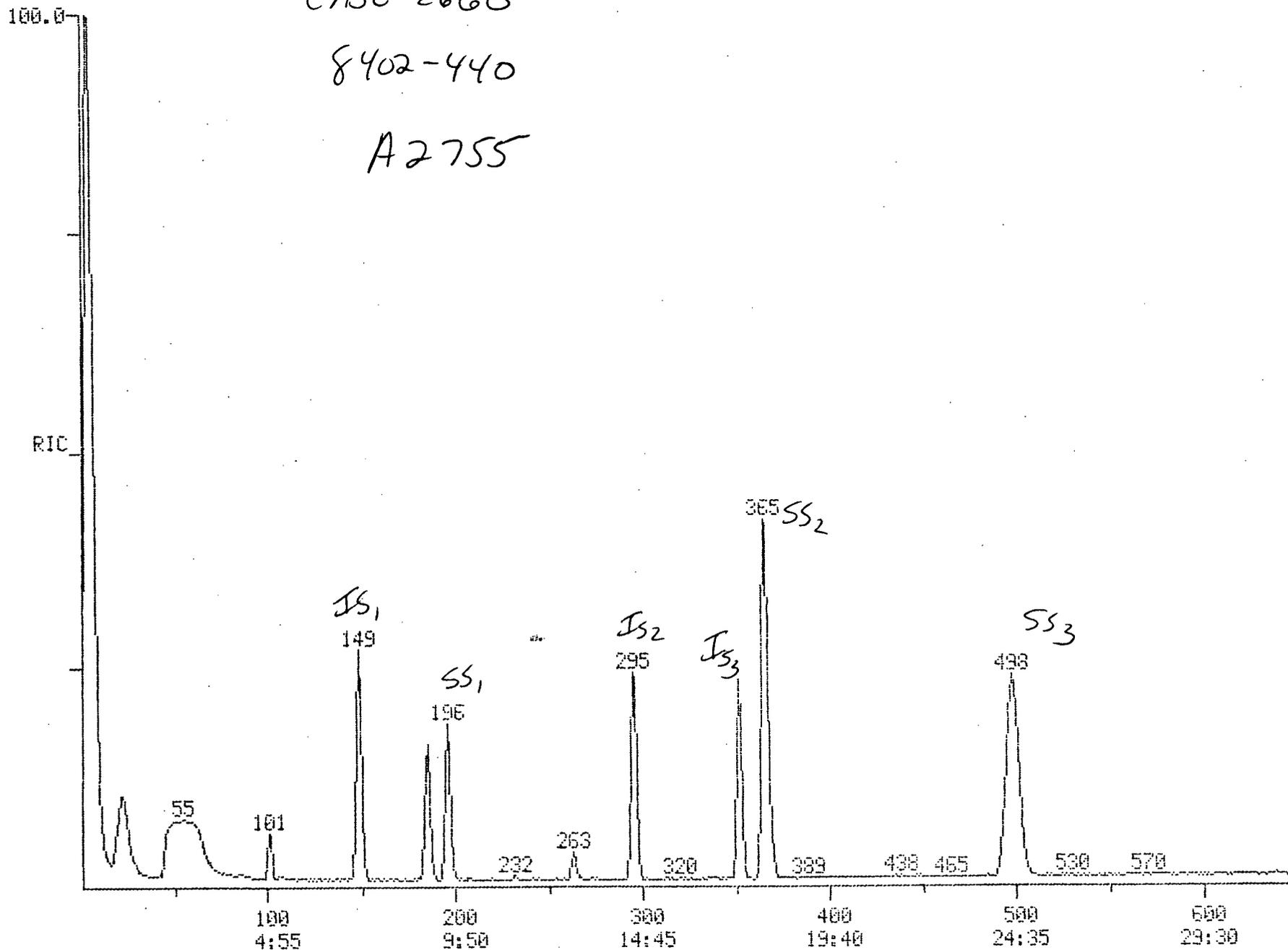
00053

CASE 2660

8402-440

A2755

161792.



A2755 (184)

SCAN
TIME

FINNIGAN ORGANICS IN WATER ANALYZER
 QUANTITATION REPORT FILE: CLPVOA33

DATA: CLPVOA33.TI
 05/05/84 21:49:00

SAMPLE:

SUBMITTED BY:

ANALYST:

AMOUNT=AREA(HGHT) * REF. AMNT / (REF. AREA(HGHT) * RESP. FACT)
 RESP. FAC. FROM LIBRARY ENTRY

- NO NAME
- 1 BROMOCHLOROMETHANE (INTERNAL STANDARD)
- 2 CHLOROMETHANE
- 3 BROMOMETHANE
- 4 VINYL CHLORIDE
- 5 CHLOROETHANE
- 6 METHYLENE CHLORIDE
- 7 CARBON DISULFIDE
- 8 1, 1 DICHLOROETHYLENE
- 9 1, 1 DICHLOROETHANE
- 10 TRANS 1, 2 DICHLOROETHYLENE
- 11 CHLOROFORM
- 12 D4-1, 2 DICHLOROETHANE (SURROGATE STANDARD)
- 13 1, 2 DICHLOROETHANE
- 14 1, 1, 1 TRICHLOROETHANE
- 15 CARBON TETRACHLORIDE
- 16 BROMODICHLOROMETHANE
- 17 2-BUTANONE (MEK)
- 18 ACETONE
- 19 ACRYLONITRILE
- 20 ACROLEIN
- 21 1-BROMO-2-CHLOROPROPANE (INTERNAL STANDARD)
- 22 1, 2 DICHLOROPROPANE
- 23 TRANS 1, 3-DICHLOROPROPENE
- 24 TRICHLOROETHYLENE
- 25 BENZENE
- 26 1, 1, 2-TRICHLOROETHANE
- 27 CIS 1, 3-DICHLOROPROPENE
- 28 DIBROMOCHLOROMETHANE
- 29 1, 4 DICHLOROBUTANE (INTERNAL STANDARD)
- 30 4-METHYL 2-PENTANONE (MIBK)
- 31 BROMOFORM
- 32 2-HEXANONE (MPK)
- 33 TETRACHLOROETHYLENE
- 34 1, 1, 2, 2 TETRACHLOROETHANE
- 35 D8-TOLUENE (SURROGATE STANDARD)
- 36 TOLUENE
- 37 CHLOROBENZENE
- 38 ETHYLBENZENE
- 39 4-BROMOFLUOROBENZENE (SURROGATE STANDARD)
- 40 STYRENE
- 41 O-XYLENE

NO	M/E	SCAN	TIME	REF	RRT	METH	AREA(HGHT)	AMOUNT	%TOT
1	130	149	7:20	1	1.000	A BB	31639.	50.000 UG/L	11.00
2	NOT FOUND								
3	NOT FOUND								

A2755 (284)

00054

NO	M/E	SCAN	TIME	REF	RRT	METH	AREA(HGHT)	AMOUNT	%TOT
4	NOT	FOUND							
5	NOT	FOUND							
6	84	101	4:58	1	0.678	A BB	5692.	2.437 UG/L	0.54 <i>BLK</i>
7	NOT	FOUND							
8	NOT	FOUND							
9	NOT	FOUND							
10	NOT	FOUND							
11	83	186	9:09	1	1.248	A BB	39651.	18.773 UG/L	4.13
12	102	196	9:38	1	1.315	A BB	4134.	88.247 PRCNT	19.41 ✓
13	NOT	FOUND							
14	NOT	FOUND							
15	NOT	FOUND							
16	NOT	FOUND							
17	NOT	FOUND							
18	43	108	5:19	1	0.725	A BB	2097.	16.256 UG/L	3.58 <i>BLK</i>
19	NOT	FOUND							
20	NOT	FOUND							
21	77	295	14:30	21	1.000	A BB	51163.	50.000 UG/L	11.00
22	NOT	FOUND							
23	NOT	FOUND							
24	130	263	12:56	21	0.892	A BB	3458.	2.652 UG/L	0.58
25	NOT	FOUND							
26	NOT	FOUND							
27	NOT	FOUND							
28	NOT	FOUND							
29	55	352	17:18	29	1.000	A BB	48435.	50.000 UG/L	11.00
30	NOT	FOUND							
31	NOT	FOUND							
32	NOT	FOUND							
33	NOT	FOUND							
34	NOT	FOUND							
35	100	365	17:57	29	1.037	A BB	74612.	79.732 PRCNT	17.54 ✓
36	92	369	18:09	29	1.048	A BB	7339.	3.666 UG/L	0.81 <i>BLK</i>
37	NOT	FOUND							
38	NOT	FOUND							
39	95	498	24:29	29	1.415	A BB	73859.	92.824 PRCNT	20.42 ✓
40	NOT	FOUND							
41	NOT	FOUND							

A2755 (3074)

00055

THE INTERNAL STANDARD AREA HAS A DIFFERENCE OF: 560 % OF THE LAST STANDARD RUN
COMPOUNDS QUANTITATED BY INTERNAL STANDARD USING PARAMETERS IN LIBRARYUX

REPORT ENTRY NO.	EXPECTED SCAN	BEST SCAN	FIT	PURITY	LIBRARY ENTRY NO.	PEAKS FOUND	PEAKS QUANT	SATURATED PEAKS
1	148	149	980	922	1	1	1	0
6	101	101	967	806	6	1	1	0
11	184	186	979	909	11	1	1	0
12	195	196	1000	790	12	1	1	0
18	108	108	759	246	18	1	1	0

THE INTERNAL STANDARD AREA HAS A DIFFERENCE OF: 7347 % OF THE LAST STANDARD RUN
COMPOUNDS QUANTITATED BY INTERNAL STANDARD USING PARAMETERS IN LIBRARYUY

REPORT ENTRY NO.	EXPECTED SCAN	BEST SCAN	FIT	PURITY	LIBRARY ENTRY NO.	PEAKS FOUND	PEAKS QUANT	SATURATED PEAKS
21	295	295	971	519	1	1	1	0
24	262	263	887	683	4	1	1	0

THE INTERNAL STANDARD AREA HAS A DIFFERENCE OF: -60 % OF THE LAST STANDARD RUN
COMPOUNDS QUANTITATED BY INTERNAL STANDARD USING PARAMETERS IN LIBRARYUZ

REPORT ENTRY NO.	EXPECTED SCAN	BEST SCAN	FIT	PURITY	LIBRARY ENTRY NO.	PEAKS FOUND	PEAKS QUANT	SATURATED PEAKS
29	355	352	989	525	1	1	1	0
35	368	365	964	865	7	1	1	0
36	372	368	954	315	8	1	1	0
COMPOUND PASSED REVERSE SEARCH BUT WAS BELOW MINIMUM AREA FOR ION QUANTIT								
37	395	368	943	14	9	1	0	0
39	506	498	987	854	11	1	1	0

NUMBER OF COMPOUNDS IDENTIFIED 12

DATA PROCESSING OF CLPVOA33 COMPLETED ON 5/05/84 22:53:56

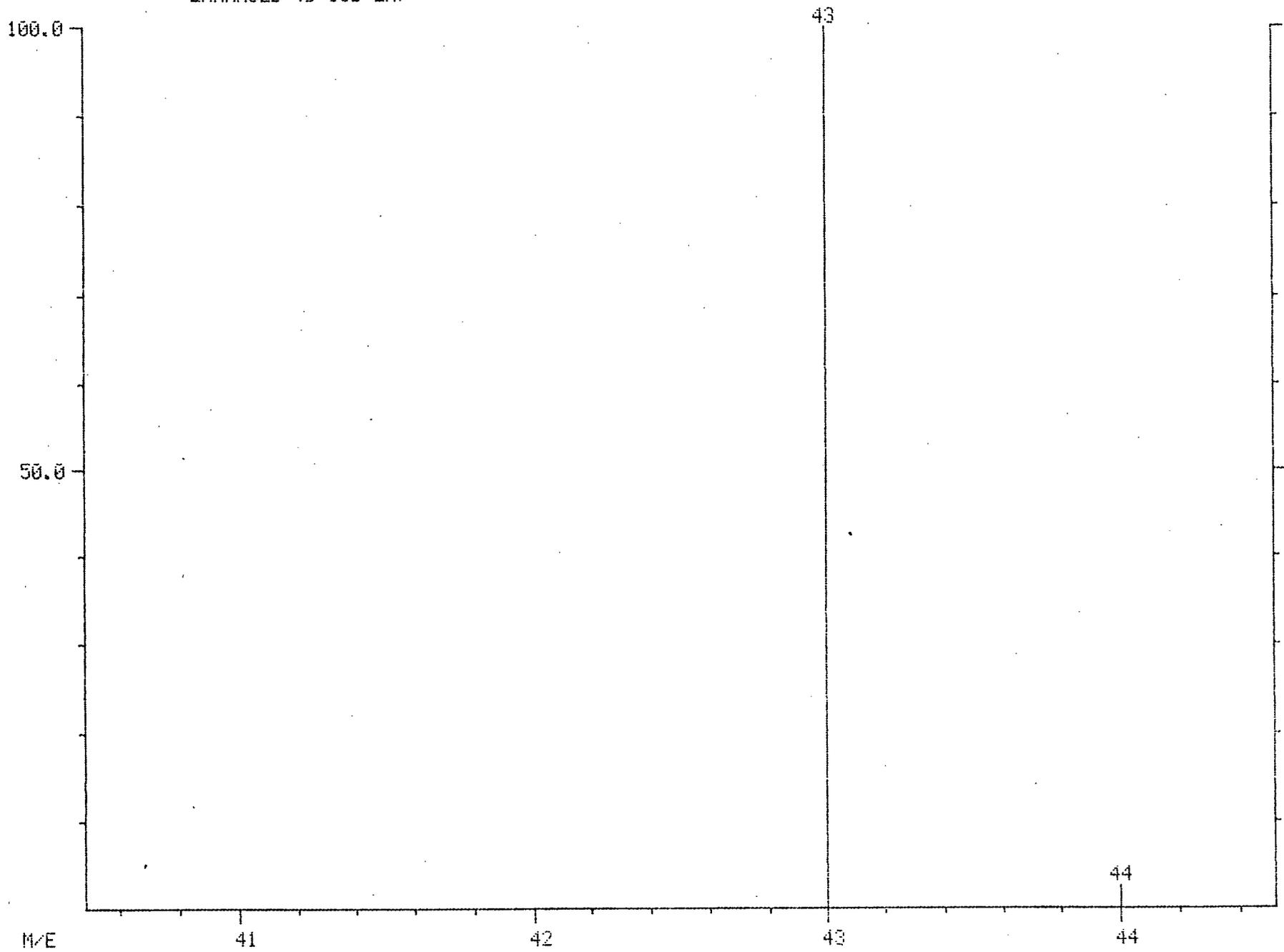
A2755 (4084)

00056

MASS SPECTRUM
05/05/84 21:49:00 + 5:19
SAMPLE:
ENHANCED (S 15B 2N)

DATA: CLP00A33 #108

BASE M/E: 43
RIC: 735.



717.
10.

A2755 - Acetone

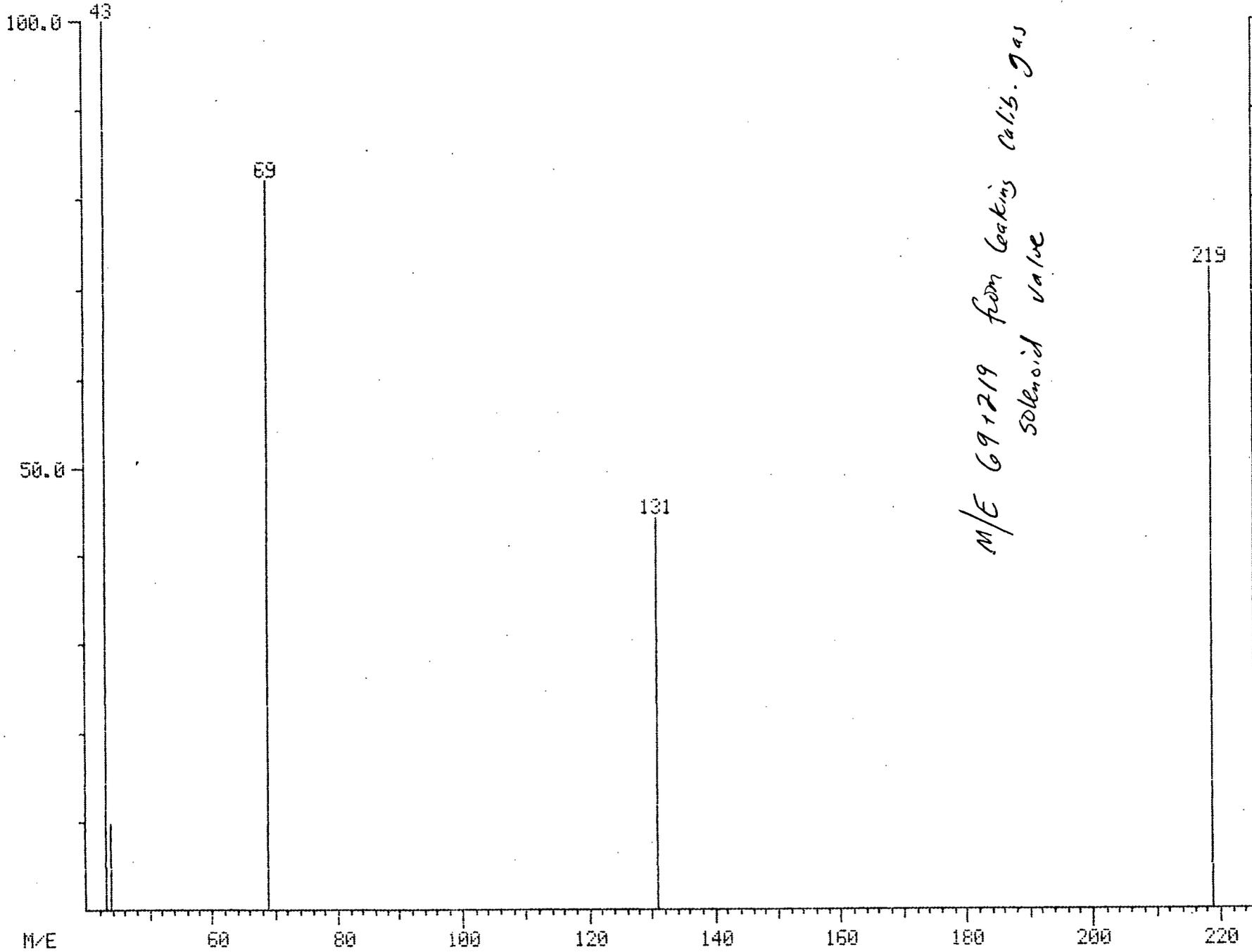
00057

8

MASS SPECTRUM
05/05/84 21:49:00 + 5:19
SAMPLE:

DATA: CLP00A33 #108

BASE M/E: 43
RIC: 2276.



*M/E 69+219 from leaking calib. gas
solenoid valve*

738.
10.

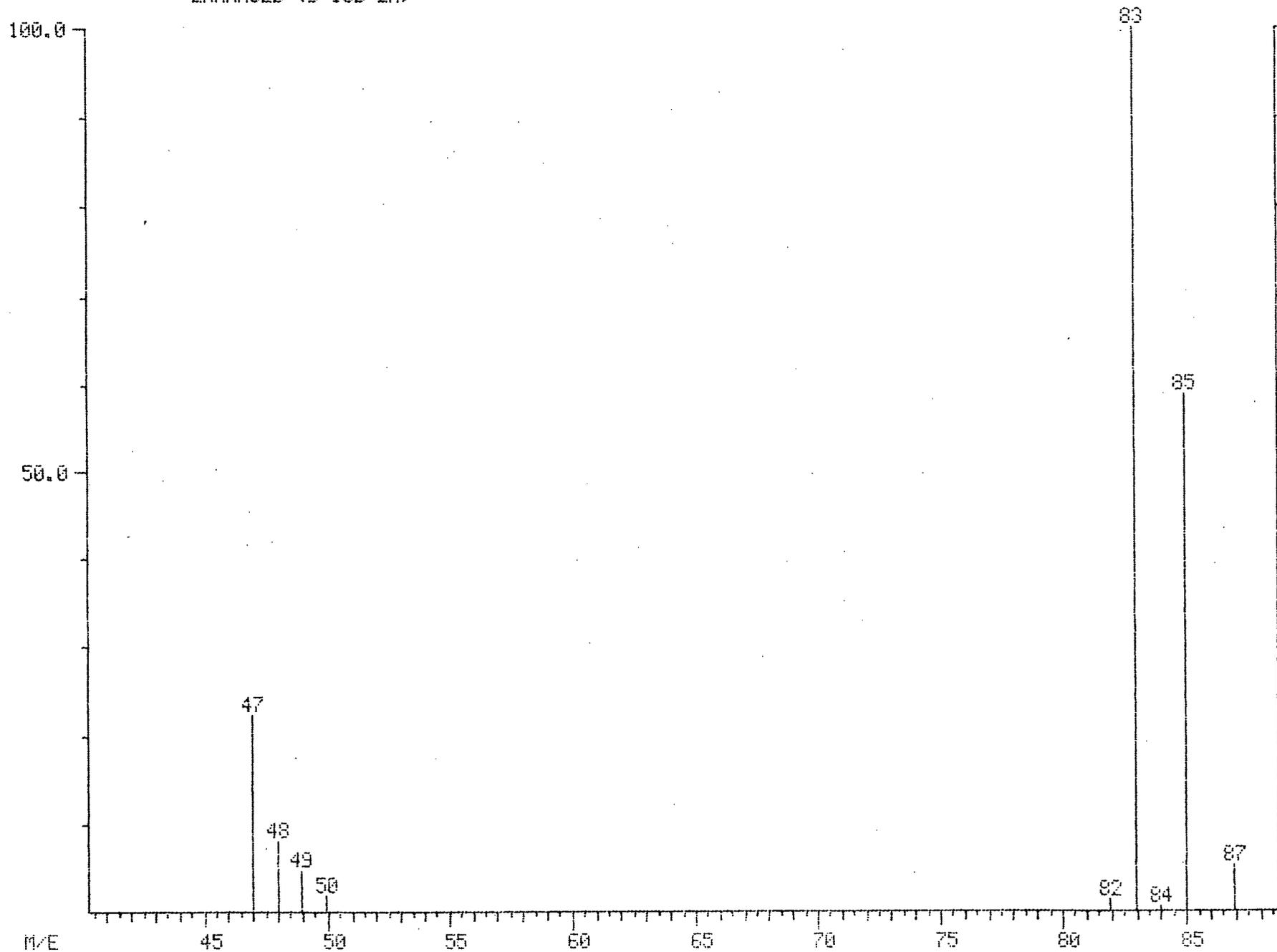
A2755- Acetone

00058

MASS SPECTRUM
05/05/84 21:49:00 + 9:09
SAMPLE:
ENHANCED (S 15B 2N)

DATA: CLP00A33 #196

BASE M/E: 83
RIC: 23392.



11536.
10.

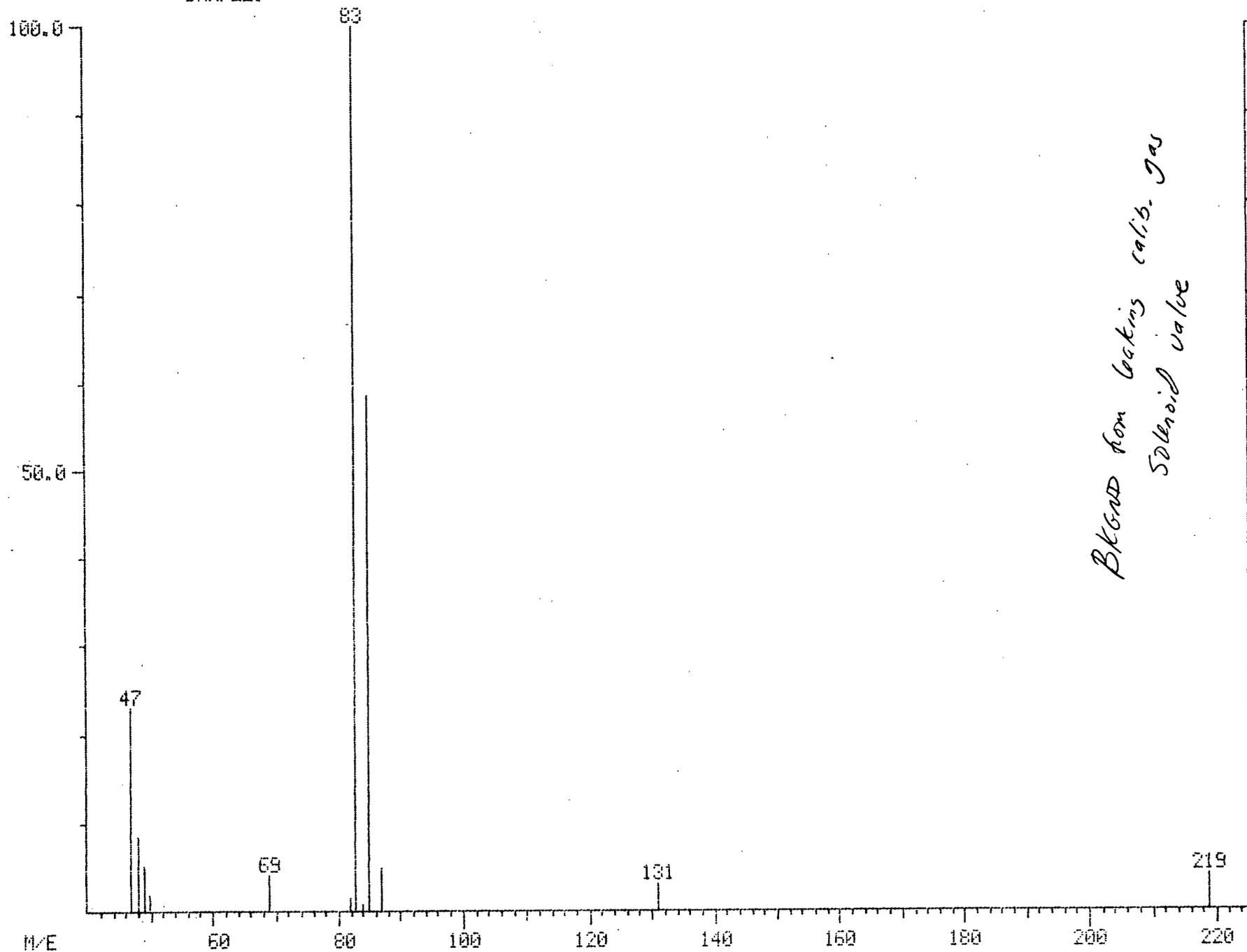
A2755- Chloroform

00059

MASS SPECTRUM
05/05/84 21:49:00 + 9:09
SAMPLE:

DATA: CLP00A33 #186

BASE M/E: 83
RIC: 26240.



*Blank from leaking calib. gas
Solenoid valve*

A2755- Chloroform

12176.
10.

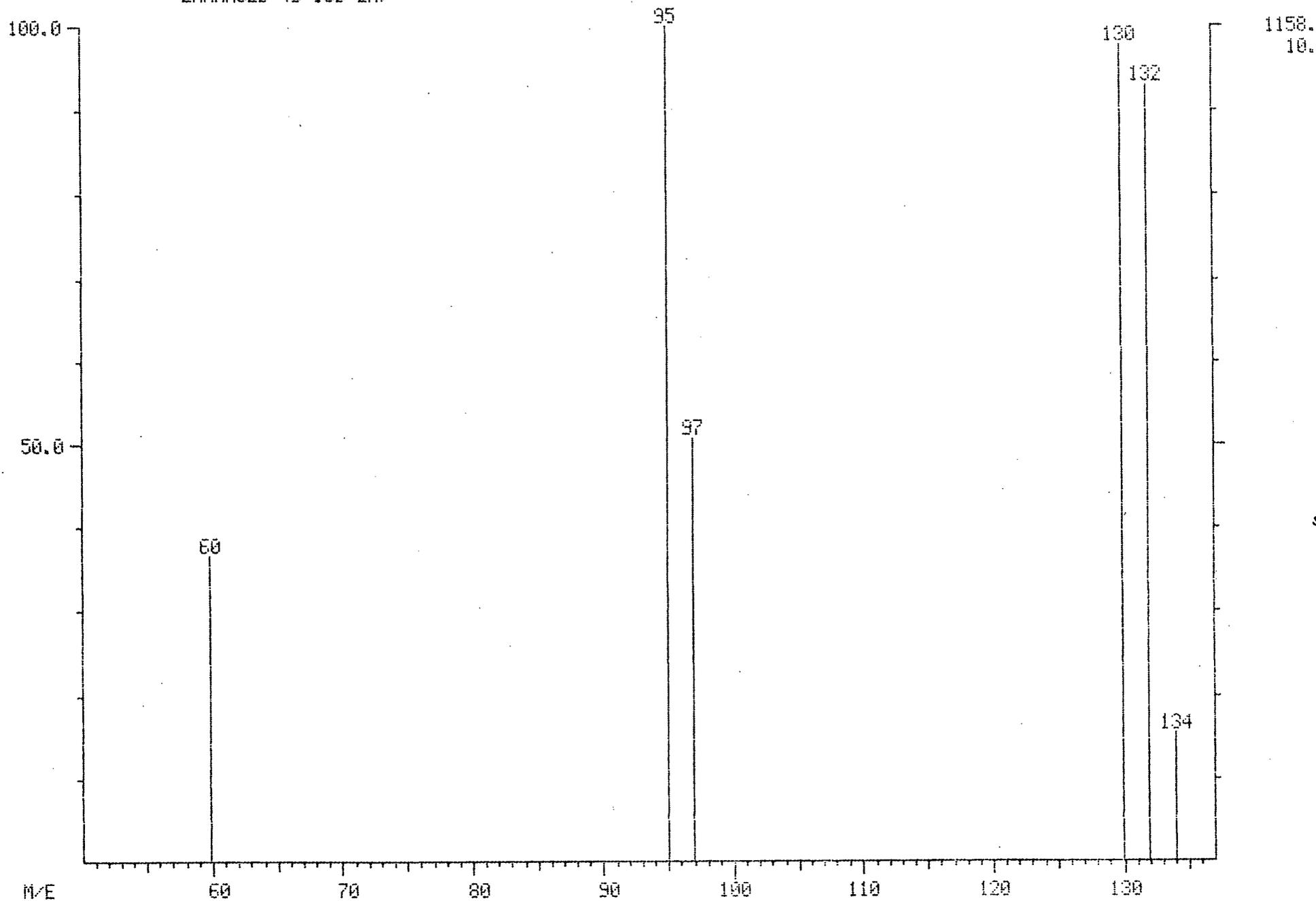
00000

MASS SPECTRUM
05/05/84 21:49:00 + 12:56
SAMPLE:
ENHANCED (S 15B 2N)

DATA: CLP00933 #263

BASE M/E: 95
RIC: 4560.

190001

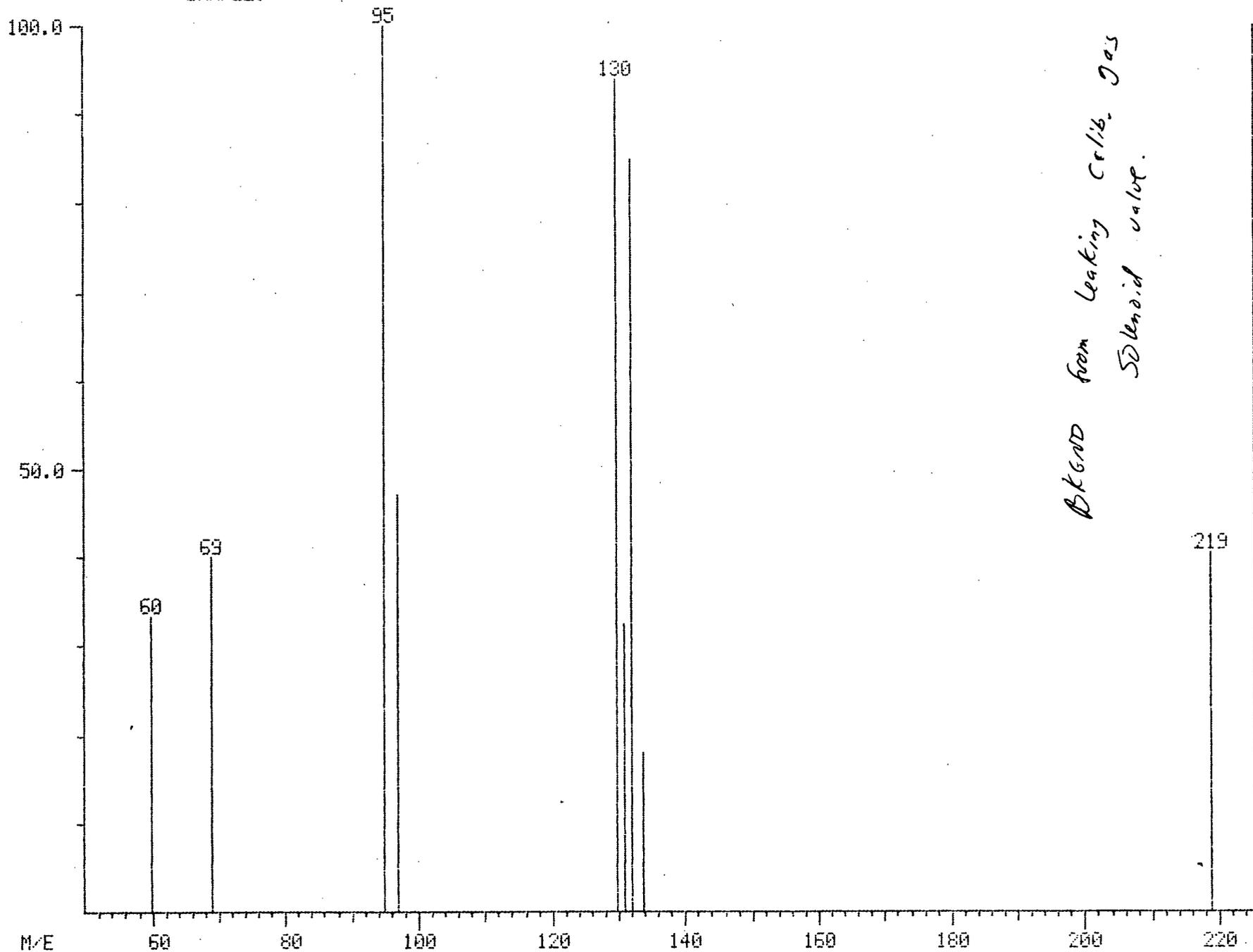


A2755- Trichloroethylene

MASS SPECTRUM
05/05/84 21:49:00 + 12:56
SAMPLE:

DATA: CLP00A33 #263

BASE M/E: 95
RIC: 6504.



*BKGD from leaking colib. Gas
Solenoid valve.*

1325.
10.

A2755- Trichloroethylene

00062

RIC
05/05/84 22:32:00
SAMPLE:

DATA: CLPVD0A34

SCANS 1 TO 650

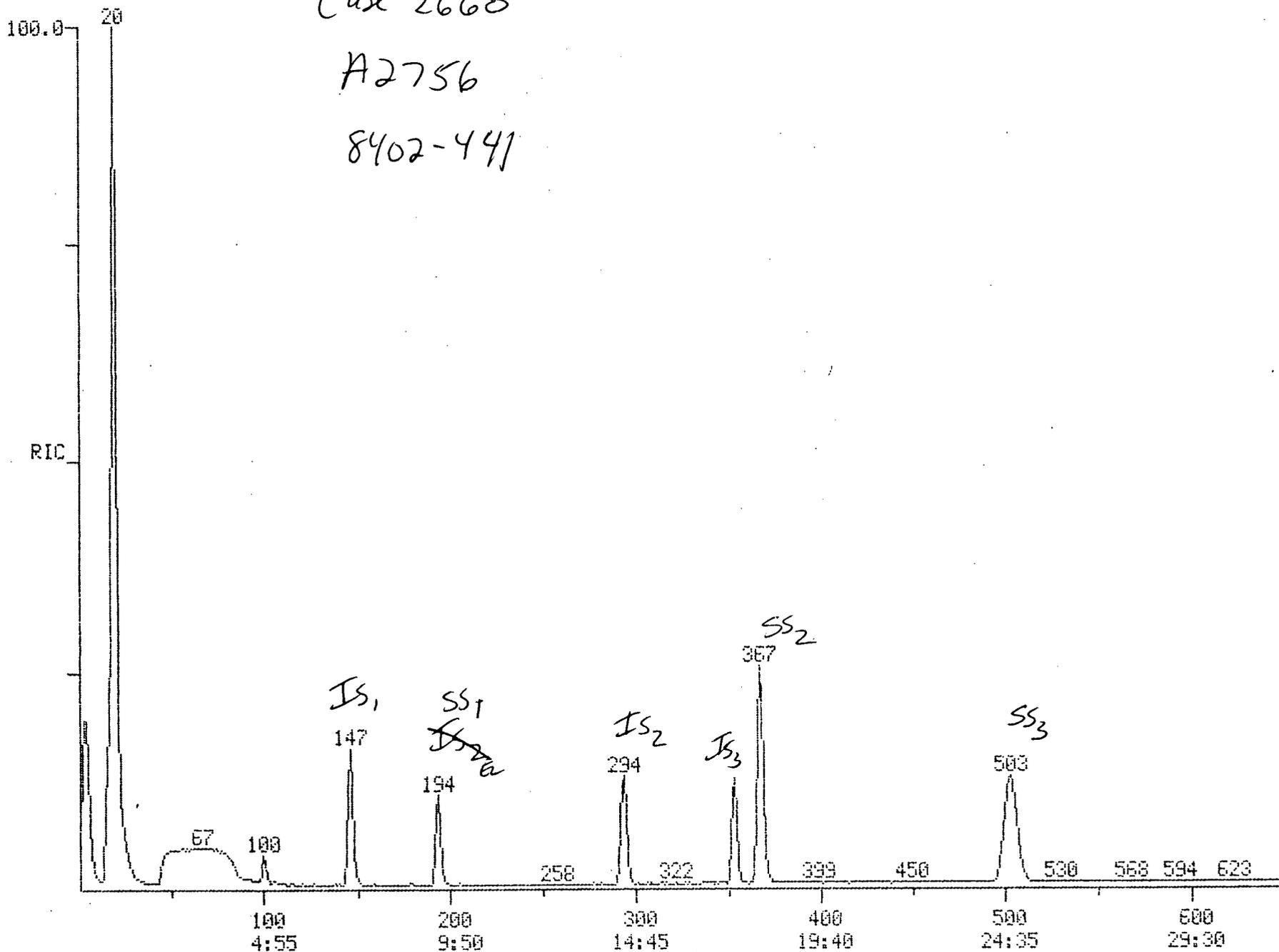
00063

Case 2660

A2756

8402-441

252160.



A2756 (1084)

SCAN
TIME

FINNIGAN ORGANICS IN WATER ANALYZER
QUANTITATION REPORT FILE: CLPVOA34

DATA: CLPVOA34.TI
05/05/84 22:32:00

SAMPLE:

SUBMITTED BY:

ANALYST:

AMOUNT=AREA(HGHT) * REF. AMNT / (REF. AREA(HGHT) * RESP. FACT)
RESP. FAC. FROM LIBRARY ENTRY

NO	NAME
1	BROMOCHLOROMETHANE (INTERNAL STANDARD)
2	CHLOROMETHANE
3	BROMOMETHANE
4	VINYL CHLORIDE
5	CHLOROETHANE
6	METHYLENE CHLORIDE
7	CARBON DISULFIDE
8	1, 1 DICHLOROETHYLENE
9	1, 1 DICHLOROETHANE
10	TRANS 1, 2 DICHLOROETHYLENE
11	CHLOROFORM
12	D4-1, 2 DICHLOROETHANE (SURROGATE STANDARD)
13	1, 2 DICHLOROETHANE
14	1, 1, 1 TRICHLOROETHANE
15	CARBON TETRACHLORIDE
16	BROMODICHLOROMETHANE
17	2-BUTANONE (MEK)
18	ACETONE
19	ACRYLONITRILE
20	ACROLEIN
21	1-BROMO-2-CHLOROPROPANE (INTERNAL STANDARD)
22	1, 2 DICHLOROPROPANE
23	TRANS 1, 3-DICHLOROPROPENE
24	TRICHLOROETHYLENE
25	BENZENE
26	1, 1, 2-TRICHLOROETHANE
27	CIS 1, 3-DICHLOROPROPENE
28	DIBROMOCHLOROMETHANE
29	1, 4 DICHLOROBUTANE (INTERNAL STANDARD)
30	4-METHYL 2-PENTANONE (MIBK)
31	BROMOFORM
32	2-HEXANONE (MPK)
33	TETRACHLOROETHYLENE
34	1, 1, 2, 2 TETRACHLOROETHANE
35	DB-TOLUENE (SURROGATE STANDARD)
36	TOLUENE
37	CHLOROBENZENE
38	ETHYLBENZENE
39	4-BROMOFLUOROBENZENE (SURROGATE STANDARD)
40	STYRENE
41	O-XYLENE

NO	M/E	SCAN	TIME	REF	RRT	METH	AREA(HGHT)	AMOUNT	%TOT
1	130	147	7:14	1	1.000	A BB	28267.	50.000 UG/L	12.09
2	NOT FOUND								
3	NOT FOUND								

A2756 (294)

00064

NO	M/E	SCAN	TIME	REF	RRT	METH	AREA(HGHT)	AMOUNT	%TOT
4		NOT FOUND							
5		NOT FOUND							
6	84	100	4:55	1	0.680	A BB	4698.	2.252 UG/L	0.54 <i>BLK</i>
7		NOT FOUND							
8		NOT FOUND							
9		NOT FOUND							
10		NOT FOUND							
11		NOT FOUND							
12	102	194	9:32	1	1.320	A BB	3570.	85.298 PRCNT	20.62 <i>✓</i>
13		NOT FOUND							
14		NOT FOUND							
15		NOT FOUND							
16		NOT FOUND							
17		NOT FOUND							
18	43	106	5:13	1	0.721	A BB	637.	5.527 UG/L	1.34 <i>BLK</i>
19		NOT FOUND							
20		NOT FOUND							
21	77	294	14:27	21	1.000	A BB	46072.	50.000 UG/L	12.09
22		NOT FOUND							
23		NOT FOUND							
24		NOT FOUND							
25		NOT FOUND							
26		NOT FOUND							
27		NOT FOUND							
28		NOT FOUND							
29	55	354	17:24	29	1.000	A BB	43147.	50.000 UG/L	12.09
30		NOT FOUND							
31		NOT FOUND							
32		NOT FOUND							
33		NOT FOUND							
34		NOT FOUND							
35	100	367	18:03	29	1.037	A BB	69347.	83.188 PRCNT	20.11 <i>✓</i>
36		NOT FOUND							
37		NOT FOUND							
38		NOT FOUND							
39	95	503	24:44	29	1.421	A BB	61986.	87.449 PRCNT	21.14 <i>✓</i>
40		NOT FOUND							
41		NOT FOUND							

A2756 (394)

00065

THE INTERNAL STANDARD AREA HAS A DIFFERENCE OF: 490 % OF THE LAST STANDARD RUN
COMPOUNDS QUANTITATED BY INTERNAL STANDARD USING PARAMETERS IN LIBRARYUX

REPORT ENTRY NO.	EXPECTED SCAN	BEST SCAN	FIT	PURITY	LIBRARY ENTRY NO.	PEAKS FOUND	PEAKS QUANT	SATURATED PEAKS
1	148	147	982	922	1	1	1	0
6	101	100	963	737	6	1	1	0
12	195	194	996	778	12	1	1	0
18	108	106	763	76	18	1	1	0

THE INTERNAL STANDARD AREA HAS A DIFFERENCE OF: 6606 % OF THE LAST STANDARD RUN
COMPOUNDS QUANTITATED BY INTERNAL STANDARD USING PARAMETERS IN LIBRARYUY

REPORT ENTRY NO.	EXPECTED SCAN	BEST SCAN	FIT	PURITY	LIBRARY ENTRY NO.	PEAKS FOUND	PEAKS QUANT	SATURATED PEAKS
21	295	294	974	493	1	1	1	0

THE INTERNAL STANDARD AREA HAS A DIFFERENCE OF: -64 % OF THE LAST STANDARD RUN
COMPOUNDS QUANTITATED BY INTERNAL STANDARD USING PARAMETERS IN LIBRARYUZ

REPORT ENTRY NO.	EXPECTED SCAN	BEST SCAN	FIT	PURITY	LIBRARY ENTRY NO.	PEAKS FOUND	PEAKS QUANT	SATURATED PEAKS
29	355	354	979	514	1	1	1	0
35	368	367	968	865	7	1	1	0
39	506	503	989	854	11	1	1	0

NUMBER OF COMPOUNDS IDENTIFIED 8
DATA PROCESSING OF CLPVOA34 COMPLETED ON 5/05/84 23:36:34

A2756 (4084)

00066

RIC
05/05/84 23:17:00
SAMPLE:

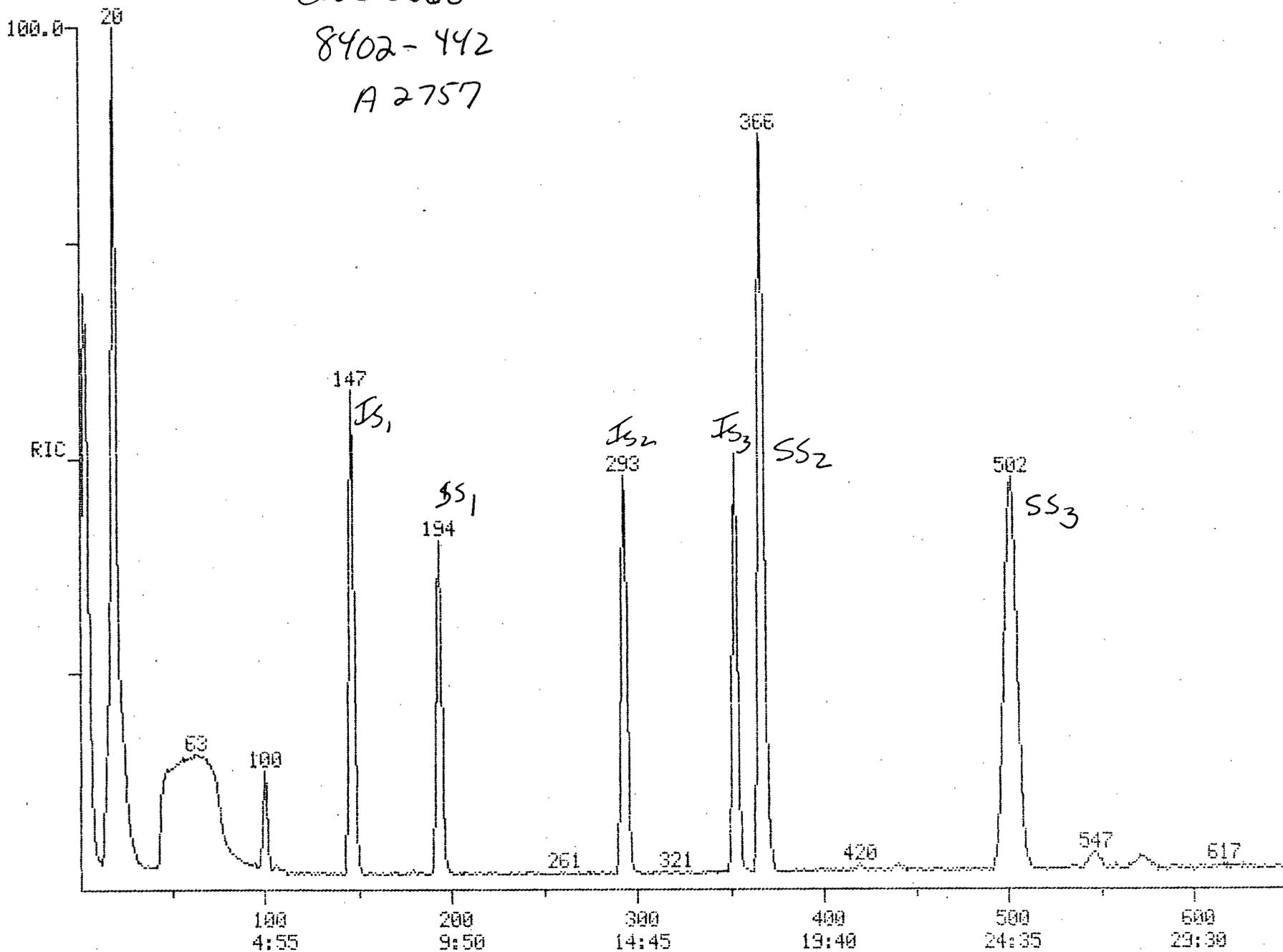
DATA: CLP00A35

SCANS 1 TO 650

00067

Case 2660
8402-442
A 2757

82048.



(15124)
A 2757 (1074)

SCAN
TIME

FINNIGAN ORGANICS IN WATER ANALYZER
 QUANTITATION REPORT FILE: CLPVOA35

DATA: CLPVOA35.TI
 05/05/84 23:17:00
 SAMPLE:
 SUBMITTED BY:

ANALYST:

AMOUNT=AREA(HGHT) * REF. AMNT/(REF. AREA(HGHT)* RESP. FACT)
 RESP. FAC. FROM LIBRARY ENTRY

- | NO | NAME |
|----|---|
| 1 | BROMOCHLOROMETHANE (INTERNAL STANDARD) |
| 2 | CHLOROMETHANE |
| 3 | BROMOMETHANE |
| 4 | VINYL CHLORIDE |
| 5 | CHLOROETHANE |
| 6 | METHYLENE CHLORIDE |
| 7 | CARBON DISULFIDE |
| 8 | 1,1 DICHLOROETHYLENE |
| 9 | 1,1 DICHLOROETHANE |
| 10 | TRANS 1,2 DICHLOROETHYLENE |
| 11 | CHLOROFORM |
| 12 | D4-1,2 DICHLOROETHANE (SURROGATE STANDARD) |
| 13 | 1,2 DICHLOROETHANE |
| 14 | 1,1,1 TRICHLOROETHANE |
| 15 | CARBON TETRACHLORIDE |
| 16 | BROMODICHLOROMETHANE |
| 17 | 2-BUTANONE (MEK) |
| 18 | ACETONE |
| 19 | ACRYLONITRILE |
| 20 | ACROLEIN |
| 21 | 1-BROMO-2-CHLOROPROPANE (INTERNAL STANDARD) |
| 22 | 1,2 DICHLOROPROPANE |
| 23 | TRANS 1,3-DICHLOROPROPENE |
| 24 | TRICHLOROETHYLENE |
| 25 | BENZENE |
| 26 | 1,1,2-TRICHLOROETHANE |
| 27 | CIS 1,3-DICHLOROPROPENE |
| 28 | DIBROMOCHLOROMETHANE |
| 29 | 1,4 DICHLOROBUTANE (INTERNAL STANDARD) |
| 30 | 4-METHYL 2-PENTANONE (MIBK) |
| 31 | BROMOFORM |
| 32 | 2-HEXANONE (MPK) |
| 33 | TETRACHLOROETHYLENE |
| 34 | 1,1,2,2 TETRACHLOROETHANE |
| 35 | D8-TOLUENE (SURROGATE STANDARD) |
| 36 | TOLUENE |
| 37 | CHLOROBENZENE |
| 38 | ETHYLBENZENE |
| 39 | 4-BROMOFLUOROBENZENE (SURROGATE STANDARD) |
| 40 | STYRENE |
| 41 | O-XYLENE |

NO	M/E	SCAN	TIME	REF	RRT	METH	AREA(HGHT)	AMOUNT	%TOT
1	130	147	7:14	1	1.000	A BB	33001.	50.000 UG/L	12.22
2	NOT FOUND								
3	NOT FOUND								

A2757 (2084)

00068

NO	M/E	SCAN	TIME	REF	RRT	METH	AREA(HGHT)	AMOUNT	%TOT
4	NOT	FOUND							
5	NOT	FOUND							
6	B4	100	4:55	1	0.680	A BE	6071.	2.492 UG/L	0.61 <i>Blk</i>
7	NOT	FOUND							
8	NOT	FOUND							
9	NOT	FOUND							
10	NOT	FOUND							
11	NOT	FOUND							
12	102	194	9:32	1	1.320	A BE	4550.	93.119 PRCNT	22.76 ✓
13	NOT	FOUND							
14	NOT	FOUND							
15	NOT	FOUND							
16	NOT	FOUND							
17	NOT	FOUND							
18	43	107	5:16	1	0.728	A BE	311.	2.311 UG/L	0.56 <i>Blk</i>
19	NOT	FOUND							
20	NOT	FOUND							
21	77	293	14:24	21	1.000	A BE	53958.	50.000 UG/L	12.22
22	NOT	FOUND							
23	NOT	FOUND							
24	NOT	FOUND							
25	NOT	FOUND							
26	NOT	FOUND							
27	NOT	FOUND							
28	NOT	FOUND							
29	55	353	17:21	29	1.000	A BE	53818.	50.000 UG/L	12.22
30	NOT	FOUND							
31	NOT	FOUND							
32	NOT	FOUND							
33	NOT	FOUND							
34	NOT	FOUND							
35	100	366	18:00	29	1.037	A BE	78622.	75.613 PRCNT	18.48 ✓
36	NOT	FOUND							
37	NOT	FOUND							
38	NOT	FOUND							
39	95	502	24:41	29	1.422	A BE	75704.	85.626 PRCNT	20.93 ✓
40	NOT	FOUND							
41	NOT	FOUND							

00069

A2757 (304)

UNKNOWN SAMPLE QUANTITATION

OWA REVERSE SEARCH STATUS REPORT

THE INTERNAL STANDARD AREA HAS A DIFFERENCE OF: 589 % OF THE LAST STANDARD RUN COMPOUNDS QUANTITATED BY INTERNAL STANDARD USING PARAMETERS IN LIBRARYUX.

REPORT ENTRY NO.	EXPECTED SCAN	BEST SCAN	FIT	PURITY	LIBRARY ENTRY NO.	PEAKS FOUND	PEAKS QUANT	SATURATED PEAKS
1	148	147	983	928	1	1	1	0
6	101	100	981	788	6	1	1	0
12	195	194	996	772	12	1	1	0
18	108	107	761	65	18	1	1	0

THE INTERNAL STANDARD AREA HAS A DIFFERENCE OF: 7754 % OF THE LAST STANDARD RUN COMPOUNDS QUANTITATED BY INTERNAL STANDARD USING PARAMETERS IN LIBRARYUY.

REPORT ENTRY NO.	EXPECTED SCAN	BEST SCAN	FIT	PURITY	LIBRARY ENTRY NO.	PEAKS FOUND	PEAKS QUANT	SATURATED PEAKS
21	295	293	973	493	1	1	1	0

THE INTERNAL STANDARD AREA HAS A DIFFERENCE OF: -55 % OF THE LAST STANDARD RUN COMPOUNDS QUANTITATED BY INTERNAL STANDARD USING PARAMETERS IN LIBRARYUZ.

REPORT ENTRY NO.	EXPECTED SCAN	BEST SCAN	FIT	PURITY	LIBRARY ENTRY NO.	PEAKS FOUND	PEAKS QUANT	SATURATED PEAKS
29	355	353	987	517	1	1	1	0
35	368	366	970	866	7	1	1	0
39	506	502	987	850	11	1	1	0

NUMBER OF COMPOUNDS IDENTIFIED 8
 DATA PROCESSING OF CLPVOA35 COMPLETED ON 5/06/84 0:18:47

A2757 (4084)

00070



Cambridge Analytical Associates

STANDARDS PACKET

00071

CASE 2660

CROSS REFERENCE TABLE FOR

VOA ANALYSIS
 BNA ANALYSIS

Cambridge Analytical

SMO #	CAA I.D.	MATRIX	SAMPLE			TUNE REFERENCE			SHIFT STANDARD			DATE OF 3 PT CURVE
			FILE	DATE	TIME	FILE	TIME	DIFF.	FILE	TIME	DIFF.	
A2749	8402-434	W	CLPVOA 23	5/5	12:04	BFBDEI	10:42	<8	CLPST 12	9:48	<8	5/5
A2750	2-435	↓	↓ 27	↓	15:13	↓	↓	<8	CLPST 12	9:49	<8	
A2751	2-436	↓	↓ 28	↓	16:00	↓	↓	<8	↓	↓	<8	
A2752	2-437	↓	↓ 29	↓	16:47	↓	↓	<8	↓	↓	<8	
A2753	2-438	↓	↓ 30	↓	17:34	↓	↓	<8	↓	↓	<8	
A2754	2-431	↓	↓ 36	↓	23:57	BFBDEI	19:46	<8	CLPSTD 13	20:09	<8	
A2755	2-440	↓	↓ 33	↓	21:49	↓	*	<8	↓	↓	<8	
A2756	2-441	↓	↓ 34	↓	22:32	↓	↓	<8	↓	↓	<8	
A2757	2-442	↓	↓	↓		↓	↓		↓	↓		

QUALITY CONTROL SAMPLES

Blank	—	W	CLPVOA 32	5/5	12:04	BFBDEI	19:46	<8	CLPSTD 13	20:09	<8	
A2754 MS	8402-434 MS	↓	↓ 24	↓	12:52	BFBDEI	10:42	<8	CLPSTD 12	9:48	<8	
A2754 MSD	2-434 MSD	↓	↓ 25	↓	13:39	↓	↓	<8	↓	9:48	<8	

00072

INITIAL CALIBRATION FILE REFERENCES

DATE	LOW LEVEL	MED. LEVEL	HIGH LEVEL
5/5	CLPSTD 10	CLPSTD 11	CLPSTD 12

ADDITIONAL COMMENTS All samples quantified against applicable 100 ppb stds.

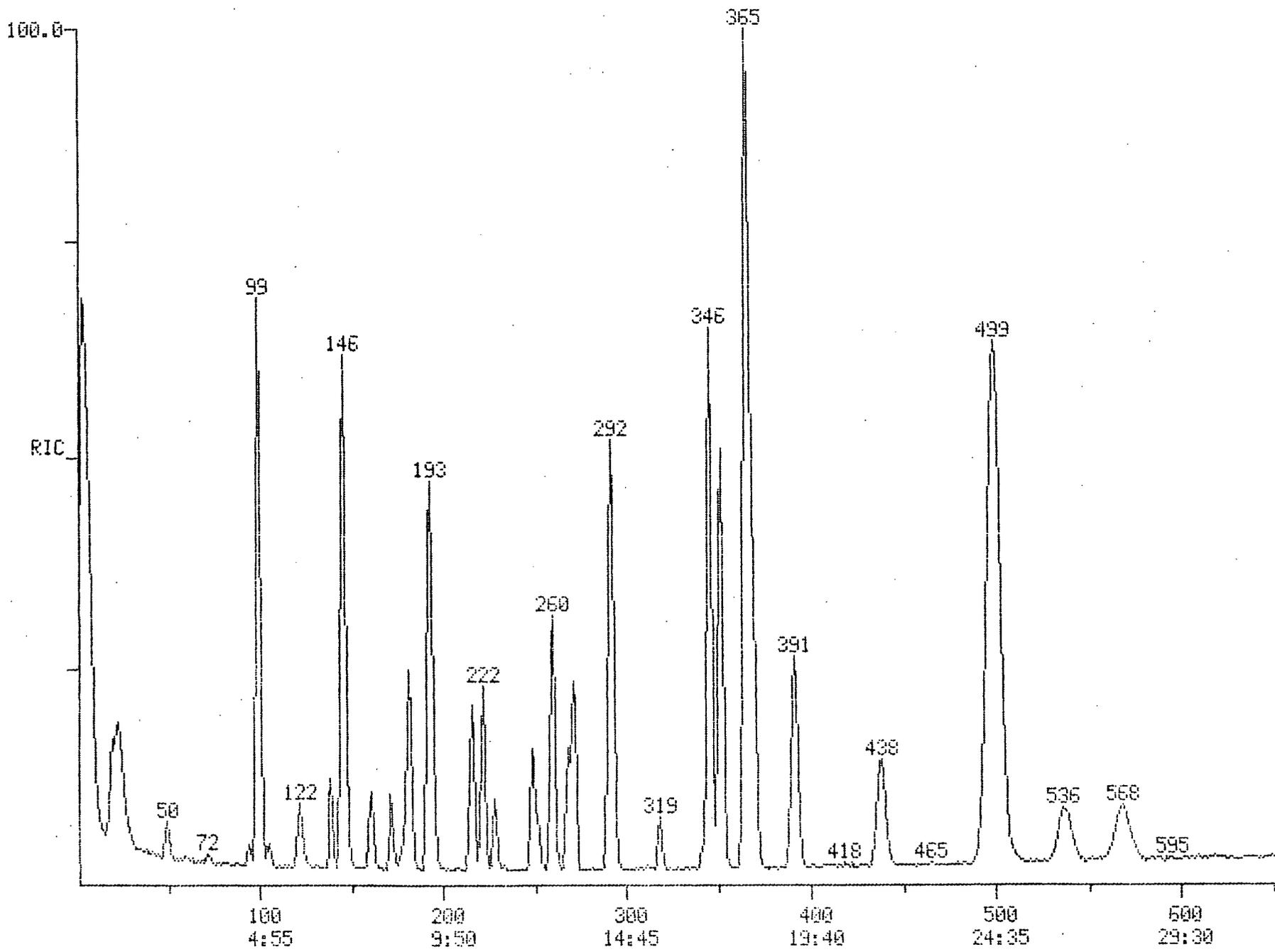
RIC
05/05/84 8:05:00
SAMPLE:

DATA: CLPSTD10

SCANS 1 TO 650

00073

97792.



20 ppb STD (1988)

SCAN
TIME

QUANTITATION REPORT FILE: CLPSTD10

DATA: CLPSTD10.TI

05/05/84 8:05:00

SAMPLE:

SUBMITTED BY:

ANALYST:

AMOUNT=AREA(HGHT) * REF. AMNT / (REF. AREA(HGHT) * RESP. FACT)
 RESP. FAC. FROM LIBRARY ENTRY

- NO NAME
- 1 BROMOCHLOROMETHANE (INTERNAL STANDARD)
- 2 CHLOROMETHANE
- 3 BROMOMETHANE
- 4 VINYL CHLORIDE
- 5 CHLOROETHANE
- 6 METHYLENE CHLORIDE
- 7 CARBON DISULFIDE
- 8 1,1 DICHLOROETHYLENE
- 9 1,1 DICHLOROETHANE
- 10 TRANS 1,2 DICHLOROETHYLENE
- 11 CHLOROFORM
- 12 D4-1,2 DICHLOROETHANE (SURROGATE STANDARD)
- 13 1,2 DICHLOROETHANE
- 14 1,1,1 TRICHLOROETHANE
- 15 CARBON TETRACHLORIDE
- 16 BROMODICHLOROMETHANE
- 17 2-BUTANONE (MEK)
- 18 ACETONE
- 19 ACRYLONITRILE
- 20 ACROLEIN
- 21 1-BROMO-2-CHLOROPROPANE (INTERNAL STANDARD)
- 22 1,2 DICHLOROPROPANE
- 23 TRANS 1,3-DICHLOROPROPENE
- 24 TRICHLOROETHYLENE
- 25 BENZENE
- 26 1,1,2-TRICHLOROETHANE
- 27 CIS 1,3-DICHLOROPROPENE
- 28 DIBROMOCHLOROMETHANE
- 29 1,4 DICHLOROETHANE (INTERNAL STANDARD)
- 30 4-METHYL 2-PENTANONE (MIBK)
- 31 BROMOFORM
- 32 2-HEXANONE (MPK)
- 33 TETRACHLOROETHYLENE
- 34 1,1,2,2 TETRACHLOROETHANE
- 35 D8-TOLUENE (SURROGATE STANDARD)
- 36 TOLUENE
- 37 CHLOROBENZENE
- 38 ETHYLBENZENE
- 39 4-BROMOFLUOROBENZENE (SURROGATE STANDARD)
- 40 STYRENE
- 41 O-XYLENE

NO	M/E	SCAN	TIME	REF	RRT	METH	AREA(HGHT)	AMOUNT	%TOT
1	130	146	7:11	1	1.000	A BB	43990.	50.000 UG/L	5.05
2	NOT FOUND								
3	94	50	2:27	1	0.342	A BB	3719.	20.000 UG/L	2.02

20 PPB VOASTD (2 of 8)

00074

NO	M/E	SCAN	TIME	REF	RRT	METH	AREA(HGHT)	AMOUNT	%TOT
4									
5									
6	84	99	4:52	1	0.678	A BB	44537.	20.000 UG/L	2.02
7	76	122	6:00	1	0.836	A BB	18999.	20.000 UG/L	2.02
8	96	139	6:50	1	0.952	A BB	6627.	20.000 UG/L	2.02
9	63	161	7:55	1	1.103	A BB	17334.	20.000 UG/L	2.02
10	96	172	8:27	1	1.178	A BB	6964.	20.000 UG/L	2.02
11	83	182	8:57	1	1.247	A BB	31175.	20.000 UG/L	2.02
12	102	193	9:29	1	1.322	A BB	6840.	100.000 PRCNT	10.10
13	62	194	9:32	1	1.329	A BB	19898.	20.000 UG/L	2.02
14	97	216	10:37	1	1.479	A BB	25842.	20.000 UG/L	2.02
15	117	222	10:55	1	1.521	A VB	21210.	20.000 UG/L	2.02
16	83	228	11:13	1	1.562	A BB	10607.	20.000 UG/L	2.02
17									
18	43	106	5:13	1	0.726	A BB	6092.	20.000 UG/L	2.02
19									
20									
21	77	292	14:21	21	1.000	A BB	63937.	50.000 UG/L	5.05
22	63	249	12:15	21	0.853	A BB	8962.	20.000 UG/L	2.02
23	75	252	12:23	21	0.863	A BB	10402.	20.000 UG/L	2.02
24	130	260	12:47	21	0.890	A BB	18884.	20.000 UG/L	2.02
25	78	268	13:11	21	0.918	A BB	25849.	20.000 UG/L	2.02
26	97	272	13:22	21	0.932	A BB	11541.	20.000 UG/L	2.02
27	75	271	13:19	21	0.928	A BB	7992.	20.000 UG/L	2.02
28	129	270	13:16	21	0.925	A BB	3379.	20.000 UG/L	2.02
29	55	352	17:18	29	1.000	A BB	63036.	50.000 UG/L	5.05
30	43	319	15:41	29	0.906	A BB	8316.	20.000 UG/L	2.02
31									
32	43	319	15:41	29	0.906	A BB	8316.	20.000 UG/L	2.02
33	164	346	17:01	29	0.983	A BB	20330.	20.000 UG/L	2.02
34	83	346	17:01	29	0.983	A BB	17860.	20.000 UG/L	2.02
35	100	365	17:57	29	1.037	A BB	100218.	100.000 PRCNT	10.10
36	92	368	18:06	29	1.045	A BB	34866.	20.000 UG/L	2.02
37	112	391	19:13	29	1.111	A BB	46184.	20.000 UG/L	2.02
38	106	438	21:32	29	1.244	A BB	12228.	20.000 UG/L	2.02
39	95	499	24:32	29	1.418	A BB	119360.	100.000 PRCNT	10.10
40									
41	106	567	27:53	29	1.611	A BB	13909.	20.000 UG/L	2.02

NO	RET(L)	RATIO	RRT(L)	RATIO	AMNT	AMNT(L)	R. FAC	R. FAC(L)	RATIO
1	7:11	1.00	1.000	1.00	50.00	50.00	1.000	1.000	1.00
2	1:55		0.260			40.00		0.095	
3	2:27	1.00	0.342	1.00	20.00	20.00	0.211	0.211	1.00
4	3:06		0.423			40.00		0.246	
5	3:44		0.510			40.00		0.184	
6	4:52	1.00	0.678	1.00	20.00	20.00	2.531	2.531	1.00
7	6:00	1.00	0.836	1.00	20.00	20.00	1.080	1.080	1.00
8	6:50	1.00	0.952	1.00	20.00	20.00	0.377	0.377	1.00
9	7:55	1.00	1.103	1.00	20.00	20.00	0.985	0.985	1.00
10	8:27	1.00	1.178	1.00	20.00	20.00	0.396	0.396	1.00
11	8:57	1.00	1.247	1.00	20.00	20.00	1.772	1.772	1.00
12	9:29	1.00	1.322	1.00	100.00	100.00	0.078	0.078	1.00
13	9:32	1.00	1.329	1.00	20.00	20.00	1.131	1.131	1.00
14	10:37	1.00	1.479	1.00	20.00	20.00	1.467	1.467	1.00
15	10:55	1.00	1.521	1.00	20.00	20.00	1.205	1.205	1.00
16	11:13	1.00	1.562	1.00	20.00	20.00	0.603	0.603	1.00

20 PPS UOASD (388) 00075

NO	RET(L)	RATIO	RRT(L)	RATIO	AMNT	AMNT(L)	R. FAC	R. FAC(L)	RATIO
17	9:35		1.309			40.00		0.230	
18	5:13	1.00	0.726	1.00	20.00	20.00	0.346	0.346	1.00
19	6:47		0.836			500.00		2.771	
20	6:29		0.943			100.00		0.014	
21	14:21	1.00	1.000	1.00	50.00	50.00	1.000	1.000	1.00
22	12:15	1.00	0.853	1.00	20.00	20.00	0.350	0.350	1.00
23	12:23	1.00	0.863	1.00	20.00	20.00	0.407	0.407	1.00
24	12:47	1.00	0.890	1.00	20.00	20.00	0.738	0.738	1.00
25	13:11	1.00	0.918	1.00	20.00	20.00	1.011	1.011	1.00
26	13:22	1.00	0.932	1.00	20.00	20.00	0.451	0.451	1.00
27	13:19	1.00	0.928	1.00	20.00	20.00	0.312	0.312	1.00
28	13:16	1.00	0.925	1.00	20.00	20.00	0.132	0.132	1.00
29	17:18	1.00	1.000	1.00	50.00	50.00	1.000	1.000	1.00
30	15:41	1.00	0.906	1.00	20.00	20.00	0.330	0.330	1.00
31	15:38		0.891			40.00		0.110	
32	15:41	1.00	0.906	1.00	20.00	20.00	0.330	0.330	1.00
33	17:01	1.00	0.983	1.00	20.00	20.00	0.806	0.806	1.00
34	17:01	1.00	0.983	1.00	20.00	20.00	0.708	0.708	1.00
35	17:57	1.00	1.037	1.00	100.00	100.00	0.795	0.795	1.00
36	18:06	1.00	1.045	1.00	20.00	20.00	1.383	1.383	1.00
37	19:13	1.00	1.111	1.00	20.00	20.00	1.832	1.832	1.00
38	21:32	1.00	1.244	1.00	20.00	20.00	0.485	0.485	1.00
39	24:32	1.00	1.418	1.00	100.00	100.00	0.947	0.947	1.00
40	26:39		1.527			40.00		1.648	
41	27:53	1.00	1.611	1.00	20.00	20.00	0.552	0.552	1.00

00076

20 ppb VDASTD (4088)

MASS CHROMATOGRAMS
05/05/84 8:05:00
SAMPLE:

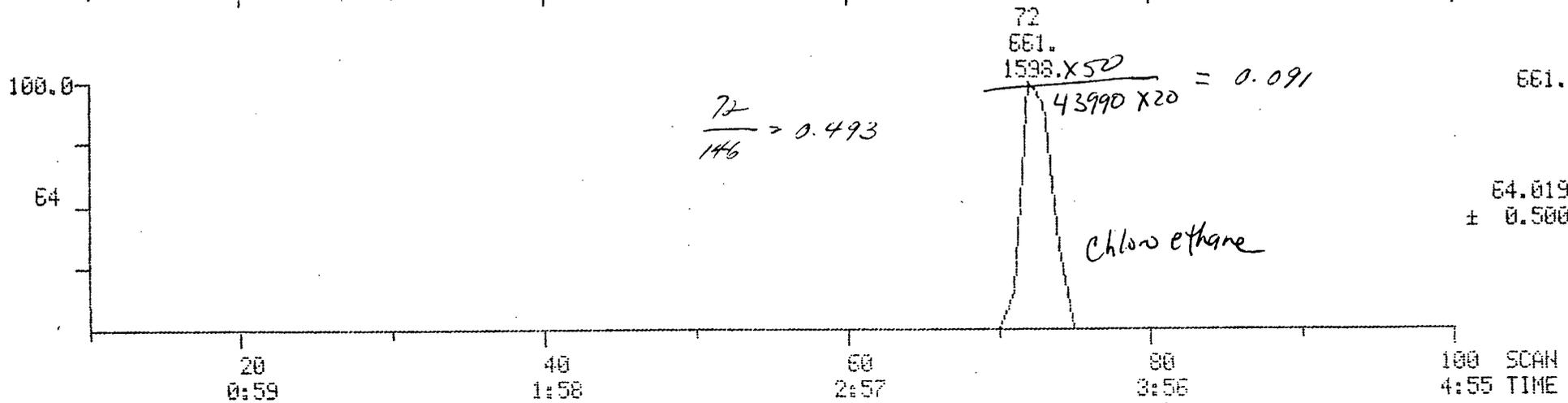
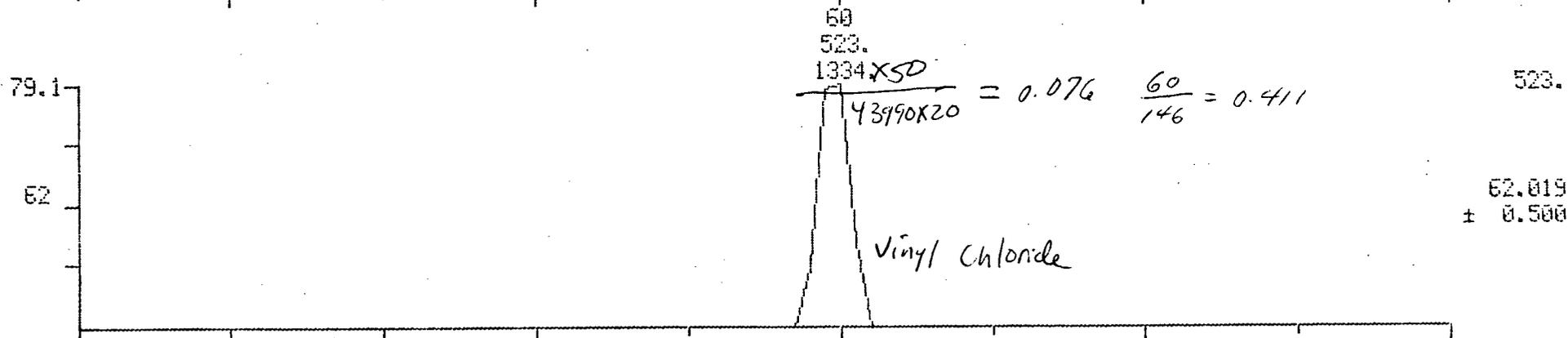
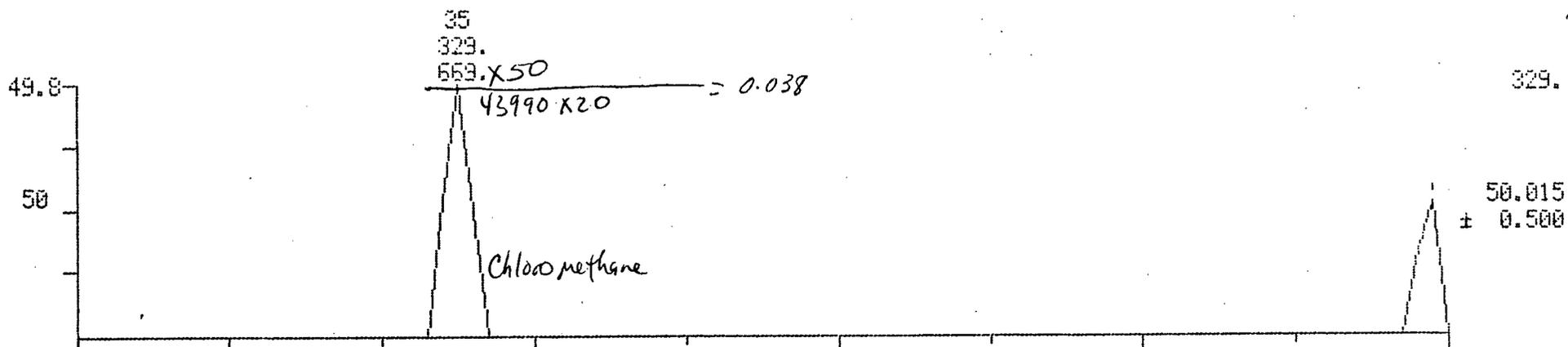
$$\frac{35}{146} =$$

DATA: CLPSTD10

SCANS 10 TO 100

00077

20 Ppb UASTD (598)

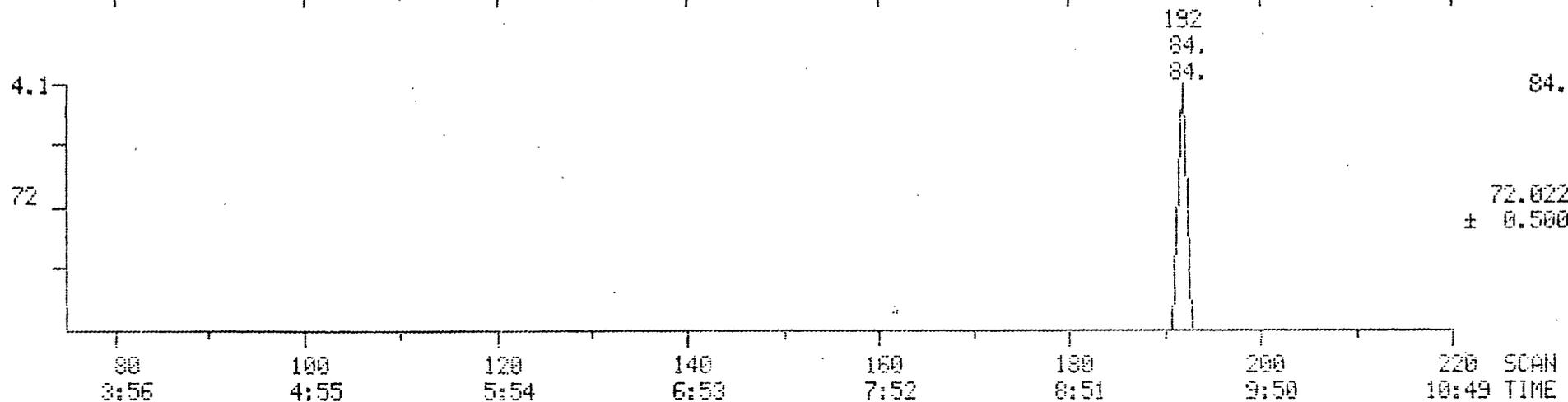
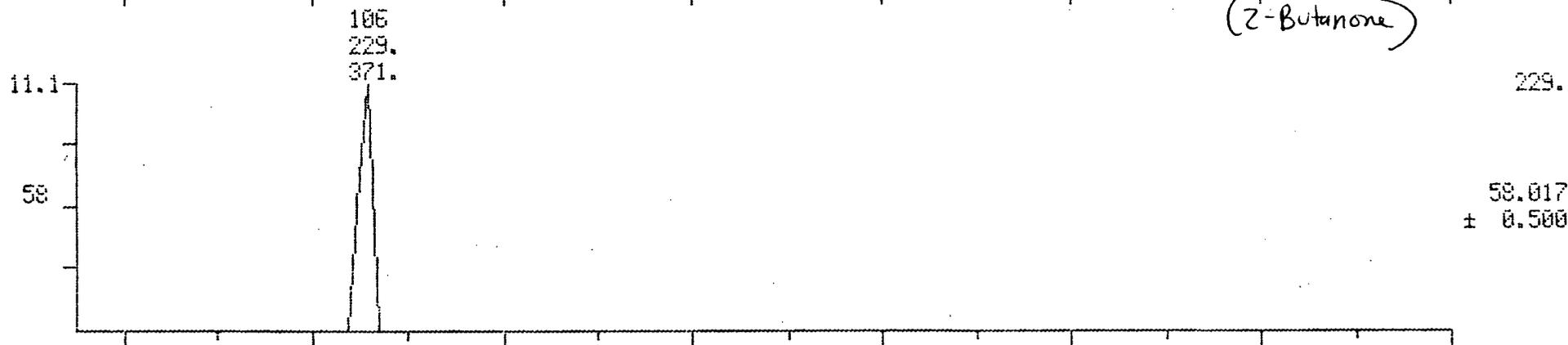
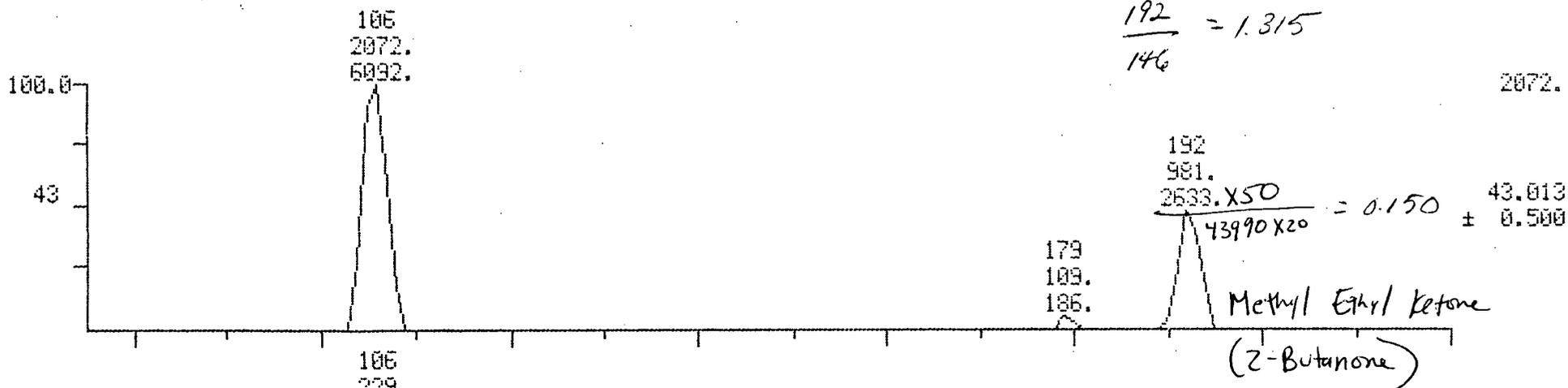


MASS CHROMATOGRAMS
05/05/84 8:05:00
SAMPLE:

DATA: CLPSTD10

SCANS 75 TO 220

00078



MASS CHROMATOGRAMS
05/05/84 8:05:00
SAMPLE:

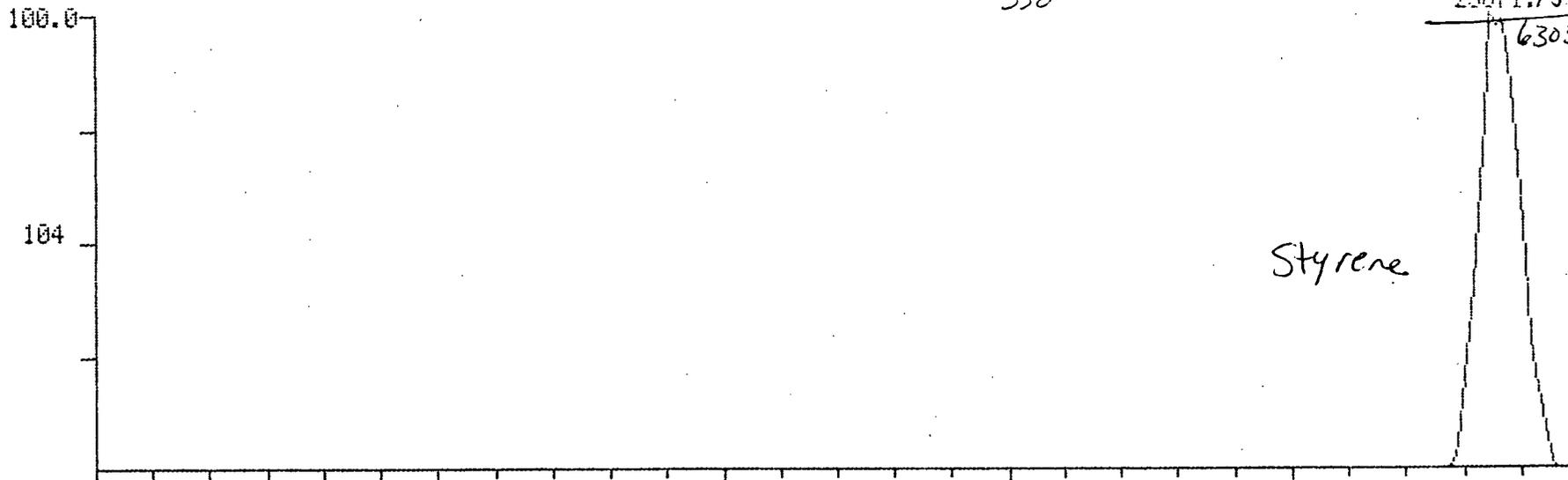
DATA: CLPSTD10

SCANS 290 TO 550

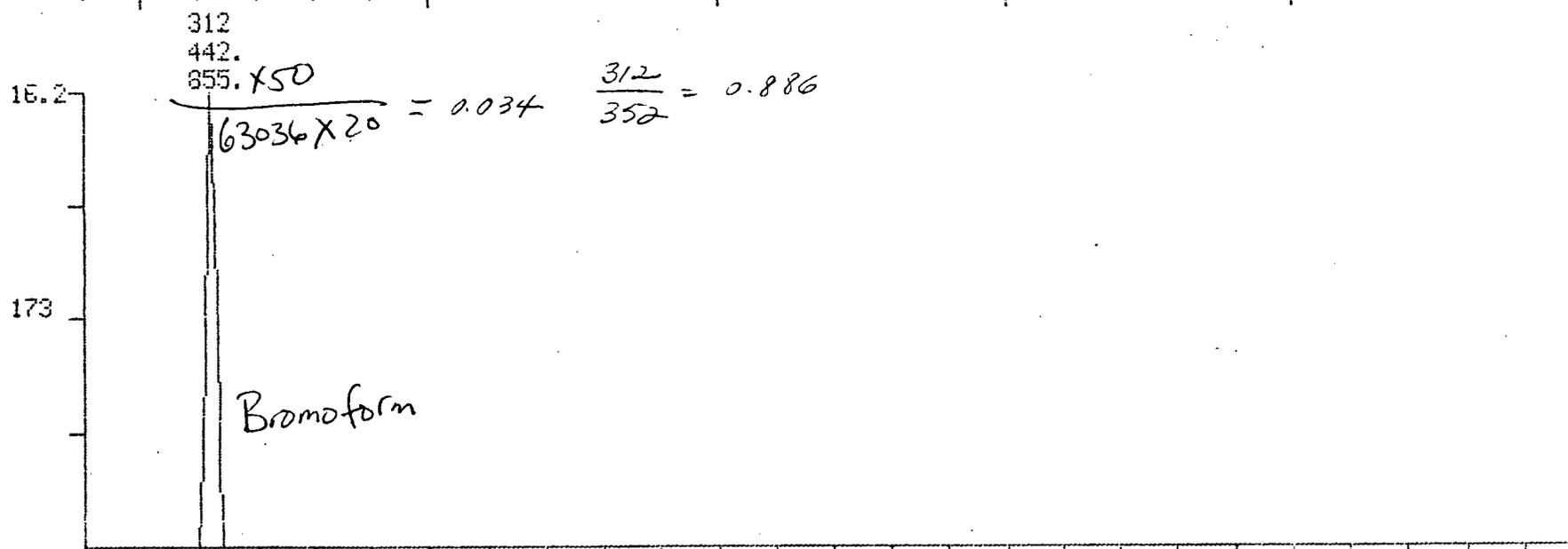
00079

DO PPS VASTD (798)

$$\frac{535}{350} = 1.520$$



$$\frac{535}{2720} = 0.1967$$
$$\frac{23571 \times 50}{63036 \times 20} = 0.939$$



$$\frac{312}{352} = 0.886$$
$$\frac{63036 \times 20}{855 \times 50} = 0.034$$

300 14:45 350 17:12 400 19:40 450 22:07 500 24:35 550 27:02 SCAN TIME

UNKNOWN SAMPLE QUANTITATION

OWA REVERSE SEARCH STATUS REPORT

THE INTERNAL STANDARD AREA HAS A DIFFERENCE OF: 62 % OF THE LAST STANDARD RUN COMPOUNDS QUANTITATED BY INTERNAL STANDARD USING PARAMETERS IN LIBRARYUX

REPORT ENTRY NO.	EXPECTED SCAN	BEST SCAN	FIT	PURITY	LIBRARY ENTRY NO.	PEAKS FOUND	PEAKS QUANT	SATURATED PEAKS
1	149	146	982	922	1	1	1	0
3	53	50	977	557	3	1	1	0
6	103	99	991	941	6	1	1	0
7	126	122	959	729	7	1	1	0
8	143	139	978	807	8	1	1	0
9	164	161	981	774	9	1	1	0
10	175	172	939	754	10	1	1	0
11	186	182	971	878	11	1	1	0
12	196	193	997	670	12	1	1	0
13	197	194	968	367	13	1	1	0
14	219	216	898	821	14	1	1	0
15	225	222	960	878	15	1	1	0
16	234	228	914	732	16	1	1	0
18	110	106	965	481	18	1	1	0

THE INTERNAL STANDARD AREA HAS A DIFFERENCE OF: 9206 % OF THE LAST STANDARD RUN COMPOUNDS QUANTITATED BY INTERNAL STANDARD USING PARAMETERS IN LIBRARYUY

REPORT ENTRY NO.	EXPECTED SCAN	BEST SCAN	FIT	PURITY	LIBRARY ENTRY NO.	PEAKS FOUND	PEAKS QUANT	SATURATED PEAKS
21	296	292	988	484	1	2	1	0
22	252	249	944	829	2	1	1	0
23	255	252	892	655	3	1	1	0
24	263	260	977	870	4	1	1	0
25	271	268	963	793	5	1	1	0
26	275	271	973	575	6	1	1	0
27	275	271	980	170	7	1	1	0
28	278	270	947	159	8	1	1	0

**

THE INTERNAL STANDARD AREA HAS A DIFFERENCE OF: -48 % OF THE LAST STANDARD RUN COMPOUNDS QUANTITATED BY INTERNAL STANDARD USING PARAMETERS IN LIBRARYUZ

REPORT ENTRY NO.	EXPECTED SCAN	BEST SCAN	FIT	PURITY	LIBRARY ENTRY NO.	PEAKS FOUND	PEAKS QUANT	SATURATED PEAKS
29	355	352	989	503	1	1	1	0
30	322	319	986	719	2	1	1	0
32	322	319	784	433	4	1	1	0
33	349	346	994	677	5	1	1	0
34	349	346	998	147	6	1	1	0
35	368	365	970	846	7	1	1	0
36	371	368	970	637	8	1	1	0
37	394	391	1000	112	9	2	1	0
38	442	438	955	786	10	1	1	0
39	503	499	989	843	11	1	1	0
40	542	537	938	628	12	1	0	0
41	573	568	871	577	13	1	1	0

**

COMPOUND PASSED REVERSE SEARCH BUT WAS BELOW MINIMUM AREA FOR ION QUANTIT

00080

NUMBER OF COMPOUNDS IDENTIFIED 34

DATA PROCESSING OF CLPSTD10 COMPLETED ON 5/05/84 8:42:14

20ppb STD(898)

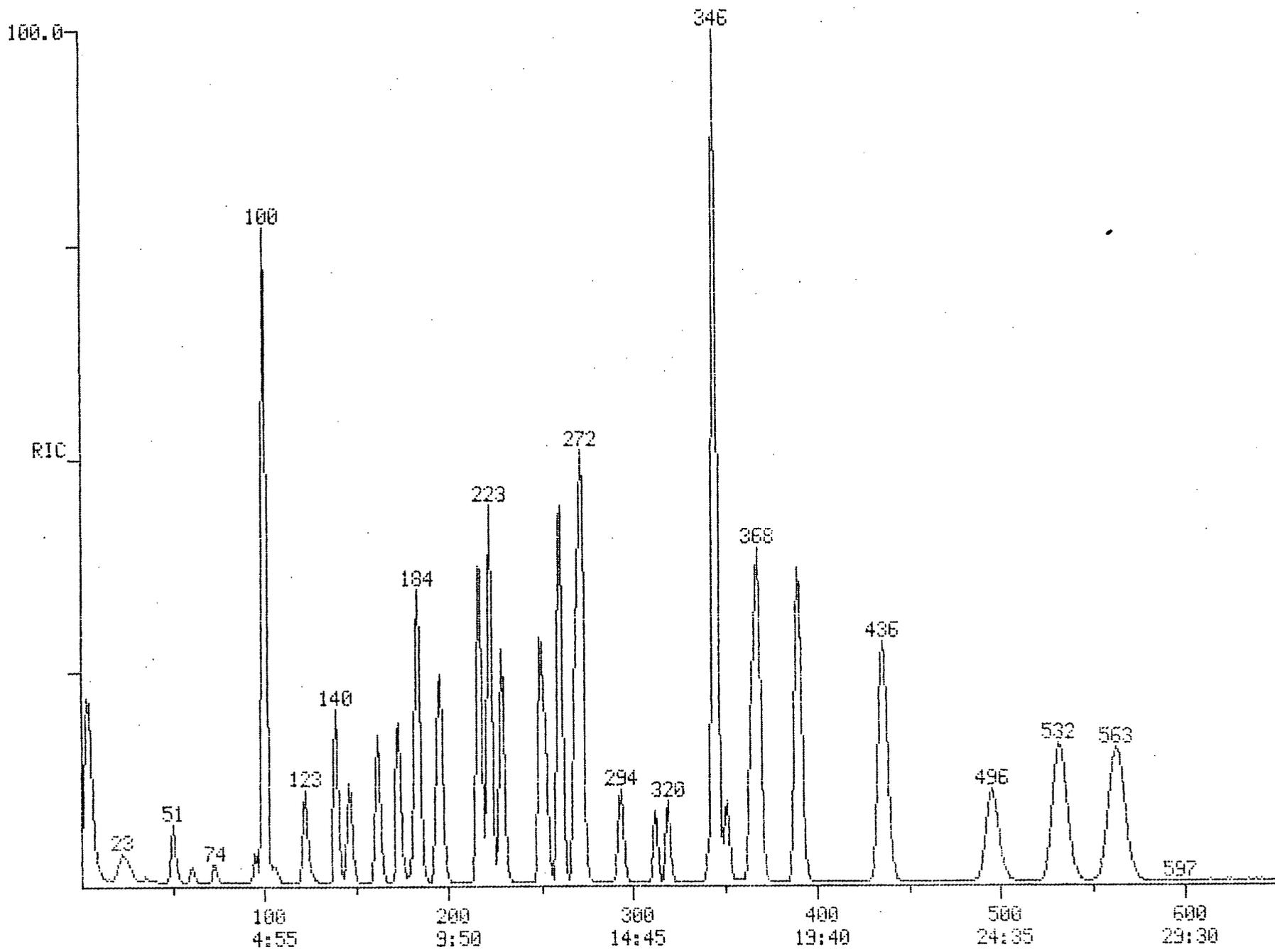
RIC
05/05/84 8:55:00
SAMPLE:

DATA: CLPSTD11

SCANS 1 TO 650

537600.

00081



100 ppb v0ASTD (1076)

SCAN
TIME

DATA: CLPSTD11.TI

05/05/84 8:55:00

SAMPLE:

SUBMITTED BY:

ANALYST:

AMOUNT=AREA(HGHT) * REF. AMNT/(REF. AREA(HGHT))* RESP. FACT)
 RESP. FAC. FROM LIBRARY ENTRY

- NO NAME
- 1 BROMOCHLOROMETHANE (INTERNAL STANDARD)
- 2 CHLOROMETHANE
- 3 BROMOMETHANE
- 4 VINYL CHLORIDE
- 5 CHLOROETHANE
- 6 METHYLENE CHLORIDE
- 7 CARBON DISULFIDE
- 8 1,1 DICHLOROETHYLENE
- 9 1,1 DICHLOROETHANE
- 10 TRANS 1,2 DICHLOROETHYLENE
- 11 CHLOROFORM
- 12 D4-1,2 DICHLOROETHANE (SURROGATE STANDARD)
- 13 1,2 DICHLOROETHANE
- 14 1,1,1 TRICHLOROETHANE
- 15 CARBON TETRACHLORIDE
- 16 BROMODICHLOROMETHANE
- 17 2-BUTANONE (MEK)
- 18 ACETONE
- 19 ACRYLONITRILE
- 20 ACROLEIN
- 21 1-BROMO-2-CHLOROPROPANE (INTERNAL STANDARD)
- 22 1,2 DICHLOROPROPANE
- 23 TRANS 1,3-DICHLOROPROPENE
- 24 TRICHLOROETHYLENE
- 25 BENZENE
- 26 1,1,2-TRICHLOROETHANE
- 27 CIS 1,3-DICHLOROPROPENE
- 28 DIBROMOCHLOROMETHANE
- 29 1,4 DICHLOROBUTANE (INTERNAL STANDARD)
- 30 4-METHYL 2-PENTANONE (MIBK)
- 31 BROMOFORM
- 32 2-HEXANONE (MPK)
- 33 TETRACHLOROETHYLENE
- 34 1,1,2,2 TETRACHLOROETHANE
- 35 D8-TOLUENE (SURROGATE STANDARD)
- 36 TOLUENE
- 37 CHLOROBENZENE
- 38 ETHYLBENZENE
- 39 4-BROMOFLUOROBENZENE (SURROGATE STANDARD)
- 40 STYRENE
- 41 O-XYLENE

NO	M/E	SCAN	TIME	REF	RRT	METH	AREA(HGHT)	AMOUNT	%TOT
1	130	147	7:14	1	1.000	A BE	47635.	50.000 UG/L	1.33
2	50	36	1:46	1	0.245	A BE	5911.	100.000 UG/L	2.67
3	94	51	2:30	1	0.347	A BE	39341.	100.000 UG/L	2.67

100 PPS UOASTD (286)

NO	M/E	SCAN	TIME	REF	RRT	METH	AREA (HGHT)	AMOUNT	%TOT
4	62	61	3:00	1	0.415	A BB	19936.	100.000 UG/L	2.67
5	64	74	3:38	1	0.503	A BB	19894.	100.000 UG/L	2.67
6	84	100	4:55	1	0.680	A BB	285229.	100.000 UG/L	2.67
7	76	123	6:03	1	0.837	A BB	160225.	100.000 UG/L	2.67
8	96	140	6:53	1	0.952	A BB	71464.	100.000 UG/L	2.67
9	63	162	7:58	1	1.102	A BB	166016.	100.000 UG/L	2.67
10	96	174	8:33	1	1.184	A BB	82237.	100.000 UG/L	2.67
11	83	184	9:03	1	1.252	A BB	282025.	100.000 UG/L	2.67
12	102	194	9:32	1	1.320	A BB	7901.	100.000 PRCNT	2.67
13	62	196	9:38	1	1.333	A BB	198495.	100.000 UG/L	2.67
14	97	217	10:40	1	1.476	A BB	256383.	100.000 UG/L	2.67
15	117	223	10:58	1	1.517	A VB	242219.	100.000 UG/L	2.67
16	83	229	11:16	1	1.558	A BB	180259.	100.000 UG/L	2.67
17	43	194	9:32	1	1.320	A BB	25753.	100.000 UG/L	2.67
18	43	107	5:16	1	0.728	A BB	18761.	100.000 UG/L	2.67
19	NOT FOUND								
20	NOT FOUND								
21	77	294	14:27	21	1.000	A BB	72889.	50.000 UG/L	1.33
22	63	250	12:17	21	0.850	A BB	95726.	100.000 UG/L	2.67
23	75	253	12:26	21	0.861	A BB	131105.	100.000 UG/L	2.67
24	130	260	12:47	21	0.884	A BB	159763.	100.000 UG/L	2.67
25	78	269	13:14	21	0.915	A BB	253244.	100.000 UG/L	2.67
26	97	273	13:25	21	0.929	A BB	103045.	100.000 UG/L	2.67
27	75	273	13:25	21	0.929	A BB	111586.	100.000 UG/L	2.67
28	129	271	13:19	21	0.922	A BB	97208.	100.000 UG/L	2.67
29	55	352	17:18	29	1.000	A BB	69876.	50.000 UG/L	1.33
30	43	320	15:44	29	0.909	A BB	72165.	100.000 UG/L	2.67
31	173	313	15:23	29	0.889	A BB	49112.	100.000 UG/L	2.67
32	43	320	15:44	29	0.909	A BB	72165.	100.000 UG/L	2.67
33	164	346	17:01	29	0.983	A BB	173795.	100.000 UG/L	2.67
34	83	346	17:01	29	0.983	A BB	152473.	100.000 UG/L	2.67
35	100	365	17:57	29	1.037	A BB	110510.	100.000 PRCNT	2.67
36	92	368	18:06	29	1.045	A BB	212806.	100.000 UG/L	2.67
37	112	390	19:10	29	1.108	A BB	347002.	100.000 UG/L	2.67
38	106	436	21:26	29	1.239	A BB	160572.	100.000 UG/L	2.67
39	95	496	24:23	29	1.409	A BB	114155.	100.000 PRCNT	2.67
40	104	532	26:09	29	1.511	A BB	285832.	100.000 UG/L	2.67
41	106	563	27:41	29	1.599	A BB	191174.	100.000 UG/L	2.67

NO	RET(L)	RATIO	RRT(L)	RATIO	AMNT	AMNT(L)	R. FAC	R. FAC(L)	RATIO
1	7:14	1.00	1.000	1.00	50.00	50.00	1.000	1.000	1.00
2	1:46	1.00	0.245	1.00	100.00	100.00	0.062	0.062	1.00
3	2:30	1.00	0.347	1.00	100.00	100.00	0.413	0.413	1.00
4	3:00	1.00	0.415	1.00	100.00	100.00	0.209	0.209	1.00
5	3:38	1.00	0.503	1.00	100.00	100.00	0.209	0.209	1.00
6	4:55	1.00	0.680	1.00	100.00	100.00	2.994	2.994	1.00
7	6:03	1.00	0.837	1.00	100.00	100.00	1.682	1.682	1.00
8	6:53	1.00	0.952	1.00	100.00	100.00	0.750	0.750	1.00
9	7:58	1.00	1.102	1.00	100.00	100.00	1.743	1.743	1.00
10	8:33	1.00	1.184	1.00	100.00	100.00	0.863	0.863	1.00
11	9:03	1.00	1.252	1.00	100.00	100.00	2.960	2.960	1.00
12	9:32	1.00	1.320	1.00	100.00	100.00	0.083	0.083	1.00
13	9:38	1.00	1.333	1.00	100.00	100.00	2.083	2.083	1.00
14	10:40	1.00	1.476	1.00	100.00	100.00	2.691	2.691	1.00
15	10:58	1.00	1.517	1.00	100.00	100.00	2.542	2.542	1.00
16	11:16	1.00	1.558	1.00	100.00	100.00	1.892	1.892	1.00

100ppb VOSTD (386)

00083

NO	RET(L)	RATIO	RRT(L)	RATIO	AMNT	AMNT(L)	R. FAC	R. FAC(L)	RATIO
17	9:32	1.00	1.320	1.00	100.00	100.00	0.270	0.270	1.00
18	5:16	1.00	0.728	1.00	100.00	100.00	0.197	0.197	1.00
19	6:47		0.836			500.00		2.771	
20	6:29		0.943			100.00		0.014	
21	14:27	1.00	1.000	1.00	50.00	50.00	1.000	1.000	1.00
22	12:17	1.00	0.850	1.00	100.00	100.00	0.657	0.657	1.00
23	12:26	1.00	0.861	1.00	100.00	100.00	0.899	0.899	1.00
24	12:47	1.00	0.884	1.00	100.00	100.00	1.096	1.096	1.00
25	13:14	1.00	0.915	1.00	100.00	100.00	1.737	1.737	1.00
26	13:25	1.00	0.929	1.00	100.00	100.00	0.707	0.707	1.00
27	13:25	1.00	0.929	1.00	100.00	100.00	0.765	0.765	1.00
28	13:19	1.00	0.922	1.00	100.00	100.00	0.667	0.667	1.00
29	17:18	1.00	1.000	1.00	50.00	50.00	1.000	1.000	1.00
30	15:44	1.00	0.909	1.00	100.00	100.00	0.516	0.516	1.00
31	15:23	1.00	0.889	1.00	100.00	100.00	0.351	0.351	1.00
32	15:44	1.00	0.909	1.00	100.00	100.00	0.516	0.516	1.00
33	17:01	1.00	0.983	1.00	100.00	100.00	1.244	1.244	1.00
34	17:01	1.00	0.983	1.00	100.00	100.00	1.091	1.091	1.00
35	17:57	1.00	1.037	1.00	100.00	100.00	0.791	0.791	1.00
36	18:06	1.00	1.045	1.00	100.00	100.00	1.523	1.523	1.00
37	19:10	1.00	1.108	1.00	100.00	100.00	2.483	2.483	1.00
38	21:26	1.00	1.239	1.00	100.00	100.00	1.149	1.149	1.00
39	24:23	1.00	1.409	1.00	100.00	100.00	0.817	0.817	1.00
40	26:09	1.00	1.511	1.00	100.00	100.00	2.045	2.045	1.00
41	27:41	1.00	1.599	1.00	100.00	100.00	1.368	1.368	1.00

100 ppb v&ASTD (496) 00084

UNKNOWN SAMPLE QUANTITATION

OWA REVERSE SEARCH STATUS REPORT

THE INTERNAL STANDARD AREA HAS A DIFFERENCE OF: 75 % OF THE LAST STANDARD RUN COMPOUNDS QUANTITATED BY INTERNAL STANDARD USING PARAMETERS IN LIBRARYUX

REPORT ENTRY NO.	EXPECTED SCAN	BEST SCAN	FIT	PURITY	LIBRARY ENTRY NO.	PEAKS FOUND	PEAKS QUANT	SATURATED PEAKS
1	146	147	981	916	1	1	1	0
2	39	36	908	404	2	1	1	0
3	50	51	984	871	3	1	1	0
4	63	61	983	722	4	1	1	0
5	76	74	990	773	5	1	1	0
6	99	100	995	953	6	1	1	0
7	122	123	986	933	7	1	1	0
8	139	140	994	909	8	1	1	0
9	161	162	994	888	9	1	1	0
10	172	174	990	933	10	1	1	0
11	182	184	991	919	11	1	1	0
12	193	194	995	411	12	1	1	0
13	194	196	980	783	13	1	1	0
14	216	217	972	922	14	1	1	0
						**		
15	222	223	984	958	15	2	1	0
16	228	229	985	907	16	1	1	0
17	195	194	980	181	17	1	1	0
						**		
18	106	107	976	549	18	2	1	0

THE INTERNAL STANDARD AREA HAS A DIFFERENCE OF: 10509 % OF THE LAST STANDARD RUN COMPOUNDS QUANTITATED BY INTERNAL STANDARD USING PARAMETERS IN LIBRARYUY

REPORT ENTRY NO.	EXPECTED SCAN	BEST SCAN	FIT	PURITY	LIBRARY ENTRY NO.	PEAKS FOUND	PEAKS QUANT	SATURATED PEAKS
						**		
21	292	294	990	439	1	2	1	0
22	249	250	979	907	2	1	1	0
23	252	253	966	771	3	1	1	0
24	260	261	990	880	4	1	1	0
25	268	269	990	774	5	1	1	0
26	272	273	992	506	6	1	1	0
27	271	272	994	203	7	1	1	0
28	270	271	1000	238	8	1	1	0

THE INTERNAL STANDARD AREA HAS A DIFFERENCE OF: -42 % OF THE LAST STANDARD RUN COMPOUNDS QUANTITATED BY INTERNAL STANDARD USING PARAMETERS IN LIBRARYUZ

REPORT ENTRY NO.	EXPECTED SCAN	BEST SCAN	FIT	PURITY	LIBRARY ENTRY NO.	PEAKS FOUND	PEAKS QUANT	SATURATED PEAKS
						**		
29	352	352	990	503	1	2	1	0
30	319	320	993	904	2	1	1	0
31	318	313	944	835	3	1	1	0
32	319	320	790	527	4	1	1	0
33	346	346	994	626	5	1	1	0
34	346	346	1000	143	6	1	1	0
35	365	365	979	718	7	1	1	0
36	368	368	990	851	8	1	1	0
						**		
37	391	390	1000	144	9	2	1	0
38	438	436	991	866	10	1	1	0
39	480	481	888	825	11	1	1	0

100 PPS
VASTD
(586)
00085

40	542	532	985	849	12	1	1	0
41	567	563	985	850	13	1	1	0

NUMBER OF COMPOUNDS IDENTIFIED 39

DATA PROCESSING OF CLPSTD11 COMPLETED ON 5/05/84 9:37:16

100PPB V0ASTD

(686) 00086

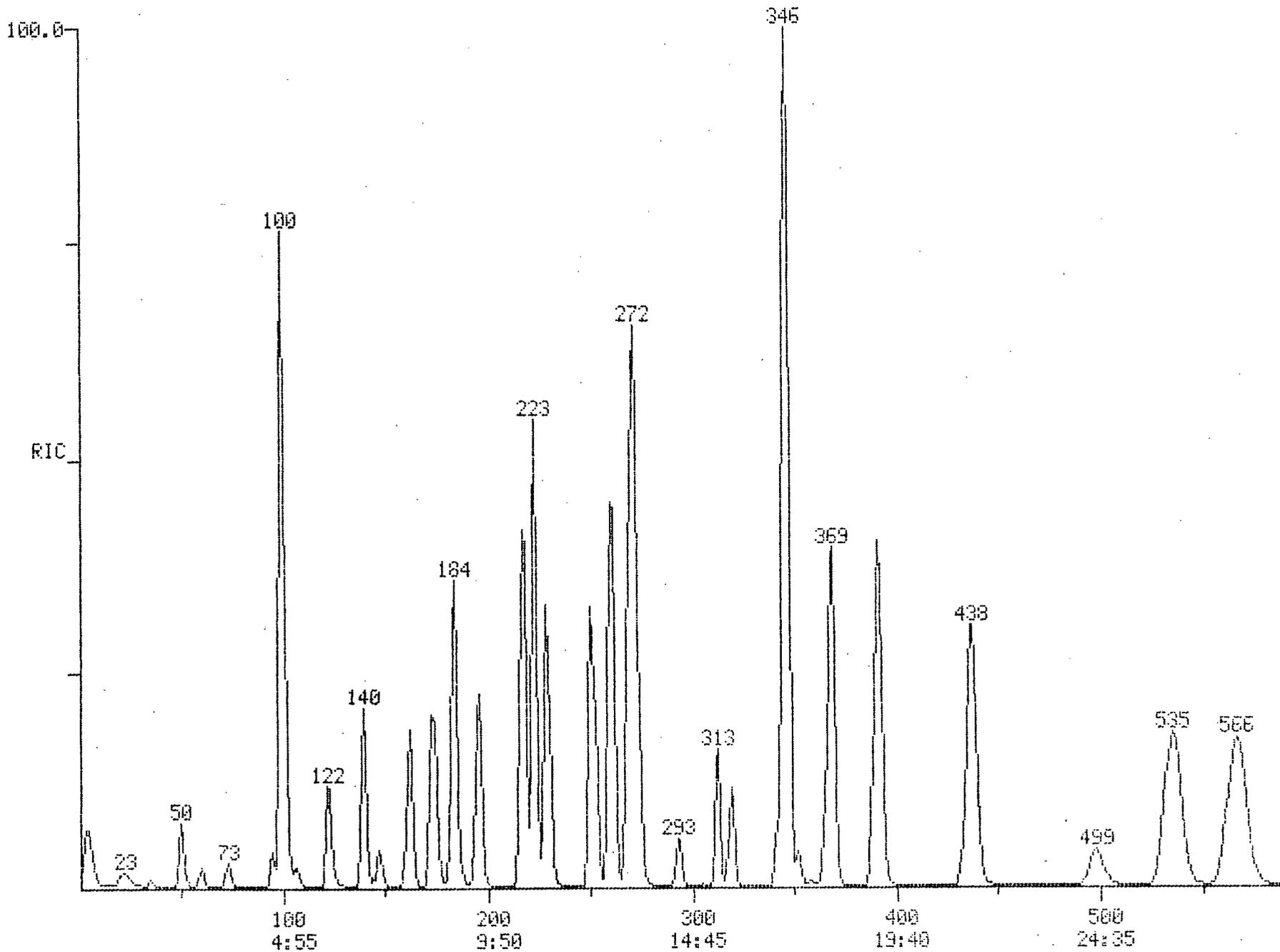
RIC
05/05/84 9:48:00
SAMPLE:

DATA: CLPSTD12

SCANS 1 TO 590

1114110.

00087



200 ppb VOA STD (1986)

SCAN
TIME

DATA: CLPSTD12.TI

05/05/84 9:48:00

SAMPLE:

SUBMITTED BY:

ANALYST:

AMOUNT=AREA(HGHT) * REF. AMNT/(REF. AREA(HGHT)* RESP. FACT)
 RESP. FAC. FROM LIBRARY ENTRY

- NO NAME
- 1 BROMOCHLOROMETHANE (INTERNAL STANDARD)
- 2 CHLOROMETHANE
- 3 BROMOMETHANE
- 4 VINYL CHLORIDE
- 5 CHLOROETHANE
- 6 METHYLENE CHLORIDE
- 7 CARBON DISULFIDE
- 8 1,1 DICHLOROETHYLENE
- 9 1,1 DICHLOROETHANE
- 10 TRANS 1,2 DICHLOROETHYLENE
- 11 CHLOROFORM
- 12 D4-1,2 DICHLOROETHANE (SURROGATE STANDARD)
- 13 1,2 DICHLOROETHANE
- 14 1,1,1 TRICHLOROETHANE
- 15 CARBON TETRACHLORIDE
- 16 BROMODICHLOROMETHANE
- 17 2-BUTANONE (MEK)
- 18 ACETONE
- 19 ACRYLONITRILE
- 20 ACRYLEIN
- 21 1-BROMO-2-CHLOROPROPANE (INTERNAL STANDARD)
- 22 1,2 DICHLOROPROPANE
- 23 TRANS 1,3-DICHLOROPROPENE
- 24 TRICHLOROETHYLENE
- 25 BENZENE
- 26 1,1,2-TRICHLOROETHANE
- 27 CIS 1,3-DICHLOROPROPENE
- 28 DIBROMOCHLOROMETHANE
- 29 1,4 DICHLOROBUTANE (INTERNAL STANDARD)
- 30 4-METHYL 2-PENTANONE (MIBK)
- 31 BROMOFORM
- 32 2-HEXANONE (MPK)
- 33 TETRACHLOROETHYLENE
- 34 1,1,2,2 TETRACHLOROETHANE
- 35 DB-TOLUENE (SURROGATE STANDARD)
- 36 TOLUENE
- 37 CHLOROBENZENE
- 38 ETHYLBENZENE
- 39 4-BROMOFLUOROBENZENE (SURROGATE STANDARD)
- 40 STYRENE
- 41 O-XYLENE

NO	M/E	SCAN	TIME	REF	RRT	METH	AREA(HGHT)	AMOUNT	%TOT
1	130	147	7:14	1	1.000	A BB	34782.	50.000 UG/L	0.49
2	50	35	1:43	1	0.238	A BB	13476.	312.228 UG/L	3.05
3	94	50	2:27	1	0.340	A BB	87051.	303.040 UG/L	2.96

200 PPS UOASTD (2 of 6)

NO	M/E	SCAN	TIME	REF	RRT	METH	AREA(HGHT)	AMOUNT	%TOT
4	62	60	2:57	1	0.408	A BB	46378.	318.600 UG/L	3.12
5	64	73	3:35	1	0.497	A BB	43885.	302.110 UG/L	2.96
6	84	100	4:55	1	0.680	A BV	592502.	284.491 UG/L	2.78
7	76	122	6:00	1	0.830	A BB	373400.	319.166 UG/L	3.12
8	96	140	6:53	1	0.952	A BB	151121.	289.607 UG/L	2.83
9	63	162	7:58	1	1.102	A BV	344775.	284.418 UG/L	2.78
10	96	173	8:30	1	1.177	A BB	179339.	298.661 UG/L	2.92
11	83	184	9:03	1	1.252	A BB	592522.	287.732 UG/L	2.82
12	102	194	9:32	1	1.320	A BB	6530.	113.189 PRCNT	1.11
13	62	196	9:38	1	1.333	A BB	417177.	287.834 UG/L	2.82
14	97	217	10:40	1	1.476	A BB	565083.	301.853 UG/L	2.95
15	117	223	10:58	1	1.517	A VB	604231.	341.638 UG/L	3.34
16	83	229	11:16	1	1.558	A BB	433463.	329.327 UG/L	3.22
17	43	194	9:32	1	1.320	A BB	64813.	344.672 UG/L	3.37
18	43	107	5:16	1	0.728	A BB	45560.	332.583 UG/L	3.25
19	NOT FOUND								
20	NOT FOUND								
21	77	293	14:24	21	1.000	A BB	61262.	50.000 UG/L	0.49
22	63	250	12:17	21	0.853	A BB	211872.	263.339 UG/L	2.58
23	75	253	12:26	21	0.863	A BB	295364.	268.046 UG/L	2.62
24	130	260	12:47	21	0.887	A BV	326481.	243.107 UG/L	2.38
25	78	269	13:14	21	0.918	A BB	572215.	268.838 UG/L	2.63
26	97	273	13:25	21	0.932	A BB	234543.	270.812 UG/L	2.65
27	75	272	13:22	21	0.928	A BB	292578.	311.963 UG/L	3.05
28	129	271	13:19	21	0.925	A BB	275681.	337.424 UG/L	3.30
29	55	352	17:18	29	1.000	A BV	58240.	50.000 UG/L	0.49
30	43	319	15:41	29	0.906	A BB	173263.	288.062 UG/L	2.82
31	173	313	15:23	29	0.889	A BB	180651.	441.326 UG/L	4.32
32	43	319	15:41	29	0.906	A BB	173263.	288.062 UG/L	2.82
33	164	346	17:01	29	0.983	A BB	358919.	247.780 UG/L	2.42
34	83	346	17:01	29	0.983	A BB	333036.	262.062 UG/L	2.56
35	100	365	17:57	29	1.037	A BB	103966.	112.875 PRCNT	1.10
36	92	369	18:09	29	1.048	A BB	438438.	247.190 UG/L	2.42
37	112	391	19:13	29	1.111	A BB	766082.	264.880 UG/L	2.59
38	106	437	21:29	29	1.241	A BB	360207.	269.147 UG/L	2.63
39	95	499	24:32	29	1.418	A BB	90102.	94.700 PRCNT	0.93
40	104	535	26:18	29	1.520	A BB	646001.	271.163 UG/L	2.65
41	106	566	27:50	29	1.608	A BB	429306.	269.428 UG/L	2.64

NO	RET(L)	RATIO	RRT(L)	RATIO	AMNT	AMNT(L)	R. FAC	R. FAC(L)	RATIO
1	7:14	1.00	1.000	1.00	50.00	50.00	1.000	1.000	1.00
2	1:46	0.97	0.245	0.97	312.23	200.00	0.097	0.062	1.56
3	2:30	0.98	0.347	0.98	303.04	200.00	0.626	0.413	1.52
4	3:00	0.98	0.415	0.98	318.60	200.00	0.333	0.209	1.59
5	3:38	0.99	0.503	0.99	302.11	200.00	0.315	0.209	1.51
6	4:55	1.00	0.680	1.00	284.49	200.00	4.259	2.994	1.42
7	6:03	0.99	0.837	0.99	319.17	200.00	2.684	1.682	1.60
8	6:53	1.00	0.952	1.00	289.61	200.00	1.086	0.750	1.45
9	7:58	1.00	1.102	1.00	284.42	200.00	2.478	1.743	1.42
10	8:33	0.99	1.184	0.99	298.66	200.00	1.289	0.863	1.49
11	9:03	1.00	1.252	1.00	287.73	200.00	4.259	2.960	1.44
12	9:32	1.00	1.320	1.00	113.19	100.00	0.094	0.083	1.13
13	9:38	1.00	1.333	1.00	287.83	200.00	2.999	2.083	1.44
14	10:40	1.00	1.476	1.00	301.85	200.00	4.062	2.691	1.51
15	10:58	1.00	1.517	1.00	341.64	200.00	4.343	2.542	1.71
16	11:16	1.00	1.558	1.00	329.33	200.00	3.116	1.892	1.65

200 PPS UOASTD (386) 00089

NO	RET(L)	RATIO	RRT(L)	RATIO	AMNT	AMNT(L)	R. FAC	R. FAC(L)	RATIO
17	9:32	1.00	1.320	1.00	344.67	200.00	0.466	0.270	1.72
18	5:16	1.00	0.728	1.00	332.58	200.00	0.327	0.197	1.66
19	6:47		0.836			500.00		2.771	
20	6:29		0.943			100.00		0.014	
21	14:27	1.00	1.000	1.00	50.00	50.00	1.000	1.000	1.00
22	12:17	1.00	0.850	1.00	263.34	200.00	0.865	0.657	1.32
23	12:26	1.00	0.861	1.00	268.05	200.00	1.205	0.899	1.34
24	12:47	1.00	0.884	1.00	243.11	200.00	1.332	1.096	1.22
25	13:14	1.00	0.915	1.00	268.84	200.00	2.335	1.737	1.34
26	13:25	1.00	0.929	1.00	270.81	200.00	0.957	0.707	1.35
27	13:25	1.00	0.929	1.00	311.96	200.00	1.194	0.765	1.56
28	13:19	1.00	0.922	1.00	337.42	200.00	1.125	0.667	1.69
29	17:18	1.00	1.000	1.00	50.00	50.00	1.000	1.000	1.00
30	15:44	1.00	0.909	1.00	288.06	200.00	0.744	0.516	1.44
31	15:23	1.00	0.889	1.00	441.33	200.00	0.775	0.351	2.21
32	15:44	1.00	0.909	1.00	288.06	200.00	0.744	0.516	1.44
33	17:01	1.00	0.983	1.00	247.78	200.00	1.541	1.244	1.24
34	17:01	1.00	0.983	1.00	262.06	200.00	1.430	1.091	1.31
35	17:57	1.00	1.037	1.00	112.87	100.00	0.893	0.791	1.13
36	18:06	1.00	1.045	1.00	247.19	200.00	1.882	1.523	1.24
37	19:10	1.00	1.108	1.00	264.88	200.00	3.288	2.483	1.32
38	21:26	1.00	1.239	1.00	269.15	200.00	1.546	1.149	1.35
39	24:23	1.01	1.409	1.01	94.70	100.00	0.774	0.817	0.95
40	26:09	1.01	1.511	1.01	271.16	200.00	2.773	2.045	1.36
41	27:41	1.01	1.599	1.01	269.43	200.00	1.843	1.368	1.35

200 PPS UOASD (486)
00090

UNKNOWN SAMPLE QUANTITATION

OWA REVERSE SEARCH STATUS REPORT

THE INTERNAL STANDARD AREA HAS A DIFFERENCE OF: 626 % OF THE LAST STANDARD RUN COMPOUNDS QUANTITATED BY INTERNAL STANDARD USING PARAMETERS IN LIBRARYUX

REPORT ENTRY NO.	EXPECTED SCAN	BEST SCAN	FIT	PURITY	LIBRARY ENTRY NO.	PEAKS FOUND	PEAKS QUANT	SATURATED PEAKS
1	147	147	982	905	1	1	1	0
2	36	35	985	667	2	1	1	0
3	51	50	989	912	3	1	1	0
4	61	60	992	853	4	1	1	0
5	74	73	995	839	5	1	1	0
6	100	100	996	951	6	1	1	0
7	123	122	989	950	7	1	1	0
8	140	140	996	905	8	1	1	0
9	162	162	996	874	9	1	1	0
10	174	173	989	936	10	1	1	0
11	184	184	993	918	11	1	1	0
12	194	194	998	207	12	1	1	0
13	196	196	971	841	13	1	1	0
14	217	217	971	918	14	1	1	0
						**		
15	223	217	989	112	15	2	1	0
16	229	229	988	903	16	1	1	0
17	194	194	992	226	17	1	1	0
						**		
18	107	107	983	648	18	2	1	0

THE INTERNAL STANDARD AREA HAS A DIFFERENCE OF: 8817 % OF THE LAST STANDARD RUN COMPOUNDS QUANTITATED BY INTERNAL STANDARD USING PARAMETERS IN LIBRARYUY

REPORT ENTRY NO.	EXPECTED SCAN	BEST SCAN	FIT	PURITY	LIBRARY ENTRY NO.	PEAKS FOUND	PEAKS QUANT	SATURATED PEAKS
						**		
21	294	293	999	383	1	2	1	0
22	250	250	980	902	2	1	1	0
23	253	253	970	773	3	1	1	0
24	260	260	993	864	4	1	1	0
25	269	269	991	689	5	1	1	0
26	273	273	996	487	6	1	1	0
27	273	272	995	213	7	1	1	0
28	271	271	997	233	8	1	1	0

THE INTERNAL STANDARD AREA HAS A DIFFERENCE OF: -52 % OF THE LAST STANDARD RUN COMPOUNDS QUANTITATED BY INTERNAL STANDARD USING PARAMETERS IN LIBRARYUZ

REPORT ENTRY NO.	EXPECTED SCAN	BEST SCAN	FIT	PURITY	LIBRARY ENTRY NO.	PEAKS FOUND	PEAKS QUANT	SATURATED PEAKS
						**		
29	352	352	988	471	1	3	1	0
30	320	319	991	903	2	1	1	0
31	313	313	966	845	3	1	1	0
32	320	319	789	515	4	1	1	0
33	346	346	995	604	5	1	1	0
34	346	346	1000	140	6	1	1	0
35	365	365	974	717	7	1	1	0
36	368	369	992	869	8	1	1	0
						**		
37	390	391	1000	149	9	2	1	0
38	436	437	988	859	10	1	1	0

200 ppb
VASTD
(586)

00091

40	532	535	991	850	12	1	1	0
41	563	566	992	847	13	1	1	0

NUMBER OF COMPOUNDS IDENTIFIED 39
DATA PROCESSING OF CLPSTD12 COMPLETED ON 5/05/84 10:26:43

200 PPS VOA STD (6036)

00092

DATA: CLPSTD13.TI

05/05/84 20:09:00

SAMPLE:

SUBMITTED BY:

ANALYST:

AMOUNT=AREA(HGHT) * REF. AMNT/(REF. AREA(HGHT)* RESP. FACT)
 RESP. FAC. FROM LIBRARY ENTRY

- NO NAME
- 1 BROMOCHLOROMETHANE (INTERNAL STANDARD)
- 2 CHLOROMETHANE
- 3 BROMOMETHANE
- 4 VINYL CHLORIDE
- 5 CHLOROETHANE
- 6 METHYLENE CHLORIDE
- 7 CARBON DISULFIDE
- 8 1,1 DICHLOROETHYLENE
- 9 1,1 DICHLOROETHANE
- 10 TRANS 1,2 DICHLOROETHYLENE
- 11 CHLOROFORM
- 12 D4-1,2 DICHLOROETHANE (SURROGATE STANDARD)
- 13 1,2 DICHLOROETHANE
- 14 1,1,1 TRICHLOROETHANE
- 15 CARBON TETRACHLORIDE
- 16 BROMODICHLOROMETHANE
- 17 2-BUTANONE (MEK)
- 18 ACETONE
- 19 ACRYLONITRILE
- 20 ACROLEIN
- 21 1-BROMO-2-CHLOROPROPANE (INTERNAL STANDARD)
- 22 1,2 DICHLOROPROPANE
- 23 TRANS 1,3-DICHLOROPROPENE
- 24 TRICHLOROETHYLENE
- 25 BENZENE
- 26 1,1,2-TRICHLOROETHANE
- 27 CIS 1,3-DICHLOROPROPENE
- 28 DIBROMOCHLOROMETHANE
- 29 1,4 DICHLOROBUTANE (INTERNAL STANDARD)
- 30 4-METHYL 2-PENTANONE (MIBK)
- 31 BROMOFORM
- 32 2-HEXANONE (MPK)
- 33 TETRACHLOROETHYLENE
- 34 1,1,2,2 TETRACHLOROETHANE
- 35 D8-TOLUENE (SURROGATE STANDARD)
- 36 TOLUENE
- 37 CHLOROBENZENE
- 38 ETHYLBENZENE
- 39 4-BROMOFLUOROBENZENE (SURROGATE STANDARD)
- 40 STYRENE
- 41 O-XYLENE

NO	M/E	SCAN	TIME	REF	RRT	METH	AREA(HGHT)	AMOUNT	%TOT
1	130	148	7:17	1	1.000	A BB	29717.	50.000 UG/L	1.17
2	NOT FOUND								
3	94	51	2:30	1	0.345	A BB	30996.	126.294 UG/L	2.97

100ppb STD (2967)

NO	M/E	SCAN	TIME	REF	RRT	METH	AREA(HGHT)	AMOUNT	%TOT
4	62	61	3:00	1	0.412	A BB	13628.	109.576 UG/L	2.57
5	64	74	3:38	1	0.500	A BB	13125.	105.754 UG/L	2.48
6	84	101	4:58	1	0.682	A BB	219339.	123.266 UG/L	2.90
7	76	124	6:06	1	0.838	A BB	119154.	119.206 UG/L	2.80
8	96	141	6:56	1	0.953	A BB	46513.	104.330 UG/L	2.45
9	63	163	8:01	1	1.101	A BB	105890.	102.241 UG/L	2.40
10	96	174	8:33	1	1.176	A BB	59400.	115.782 UG/L	2.72
11	83	184	9:03	1	1.243	A BB	198384.	112.756 UG/L	2.65
12	102	195	9:35	1	1.318	A BB	4400.	89.267 PRCNT	2.10
13	62	197	9:41	1	1.331	A BB	134243.	108.409 UG/L	2.55
14	97	219	10:46	1	1.480	A BB	197590.	123.537 UG/L	2.90
15	117	225	11:04	1	1.520	A VB	203472.	134.654 UG/L	3.16
16	83	231	11:21	1	1.561	A BB	140784.	125.192 UG/L	2.94
17	43	195	9:35	1	1.318	A BB	13661.	85.031 UG/L	2.00
18	43	108	5:19	1	0.730	A BB	12116.	103.520 UG/L	2.43
19	NOT FOUND								
20	NOT FOUND								
21	77	295	14:30	21	1.000	A BB	45246.	50.000 UG/L	1.17
22	63	252	12:23	21	0.854	A BB	59574.	100.256 UG/L	2.36
23	75	255	12:32	21	0.864	A BB	95954.	117.903 UG/L	2.77
24	130	262	12:53	21	0.888	A BB	115329.	116.276 UG/L	2.73
25	78	271	13:19	21	0.919	A BB	181225.	115.282 UG/L	2.71
26	97	274	13:28	21	0.929	A BB	73871.	115.486 UG/L	2.71
27	75	274	13:28	21	0.929	A BB	89032.	128.534 UG/L	3.02
28	129	273	13:25	21	0.925	A BB	100236.	166.113 UG/L	3.90
29	55	355	17:27	29	1.000	A BB	37984.	50.000 UG/L	1.17
30	43	322	15:50	29	0.907	A BB	41377.	105.477 UG/L	2.48
31	173	315	15:29	29	0.887	A BB	59007.	221.026 UG/L	5.19
32	43	322	15:50	29	0.907	A BB	41377.	105.477 UG/L	2.48
33	164	349	17:10	29	0.983	A BB	137904.	145.971 UG/L	3.43
34	83	349	17:10	29	0.983	A BB	103051.	125.298 UG/L	2.94
35	100	368	18:06	29	1.037	A BB	73387.	122.164 PRCNT	2.87
36	92	372	18:17	29	1.048	A BB	156979.	135.702 UG/L	3.19
37	112	395	19:25	29	1.113	A BB	255284.	135.338 UG/L	3.18
38	106	443	21:47	29	1.248	A BB	110822.	126.965 UG/L	2.98
39	95	506	24:53	29	1.425	A BB	62400.	100.559 PRCNT	2.36
40	104	544	26:45	29	1.532	A BB	207020.	133.238 UG/L	3.13
41	NOT FOUND								

NO	RET(L)	RATIO	RRT(L)	RATIO	AMNT	AMNT(D)	R. FAC	R. FAC(L)	RATIO
1	7:14	1.01	1.000	1.00	50.00	50.00	1.000	1.000	1.00
2	1:46		0.245			200.00		0.062	
3	2:30	1.00	0.347	0.99	126.29	100.00	0.522	0.413	1.26
4	3:00	1.00	0.415	0.99	109.58	100.00	0.229	0.209	1.10
5	3:38	1.00	0.503	0.99	105.75	100.00	0.221	0.209	1.06
6	4:55	1.01	0.680	1.00	123.27	100.00	3.690	2.994	1.23
7	6:03	1.01	0.837	1.00	119.21	100.00	2.005	1.682	1.19
8	6:53	1.01	0.952	1.00	104.33	100.00	0.783	0.750	1.04
9	7:58	1.01	1.102	1.00	102.24	100.00	1.782	1.743	1.02
10	8:33	1.00	1.184	0.99	115.78	100.00	0.999	0.863	1.16
11	9:03	1.00	1.252	0.99	112.76	100.00	3.338	2.960	1.13
12	9:32	1.01	1.320	1.00	89.27	100.00	0.074	0.083	0.89
13	9:38	1.01	1.333	1.00	108.41	100.00	2.259	2.083	1.08
14	10:40	1.01	1.476	1.00	123.54	100.00	3.325	2.691	1.24
15	10:58	1.01	1.517	1.00	134.65	100.00	3.423	2.542	1.35
16	11:16	1.01	1.558	1.00	125.19	100.00	2.369	1.892	1.25

100 ppb STD (387)

00095

NO	RET(L)	RATIO	RRT(L)	RATIO	AMNT	AMNT(L)	R. FAC	R. FAC(L)	RATIO
17	9:32	1.01	1.320	1.00	85.03	100.00	0.230	0.270	0.85
18	5:16	1.01	0.728	1.00	103.52	100.00	0.204	0.197	1.04
19	6:47		0.836			500.00		2.771	
20	6:29		0.743			100.00		0.014	
21	14:27	1.00	1.000	1.00	50.00	50.00	1.000	1.000	1.00
22	12:17	1.01	0.850	1.00	100.26	100.00	0.658	0.657	1.00
23	12:26	1.01	0.861	1.00	117.90	100.00	1.060	0.899	1.18
24	12:47	1.01	0.884	1.00	116.28	100.00	1.274	1.096	1.16
25	13:14	1.01	0.915	1.00	115.28	100.00	2.003	1.737	1.15
26	13:25	1.00	0.929	1.00	115.49	100.00	0.816	0.707	1.15
27	13:25	1.00	0.929	1.00	128.53	100.00	0.984	0.765	1.29
28	13:19	1.01	0.922	1.00	166.11	100.00	1.108	0.667	1.66
29	17:18	1.01	1.000	1.00	50.00	50.00	1.000	1.000	1.00
30	15:44	1.01	0.909	1.00	105.48	100.00	0.545	0.516	1.05
31	15:23	1.01	0.889	1.00	221.03	100.00	0.777	0.351	2.21
32	15:44	1.01	0.909	1.00	105.48	100.00	0.545	0.516	1.05
33	17:01	1.01	0.983	1.00	145.97	100.00	1.815	1.244	1.46
34	17:01	1.01	0.983	1.00	125.30	100.00	1.367	1.091	1.25
35	17:57	1.01	1.037	1.00	122.16	100.00	0.966	0.791	1.22
36	18:06	1.01	1.045	1.00	135.70	100.00	2.066	1.523	1.36
37	19:10	1.01	1.108	1.00	135.34	100.00	3.360	2.483	1.35
38	21:26	1.02	1.239	1.01	126.96	100.00	1.459	1.149	1.27
39	24:23	1.02	1.409	1.01	100.56	100.00	0.821	0.817	1.01
40	26:09	1.02	1.511	1.01	133.24	100.00	2.725	2.045	1.33
41	27:41		1.599			200.00		1.368	

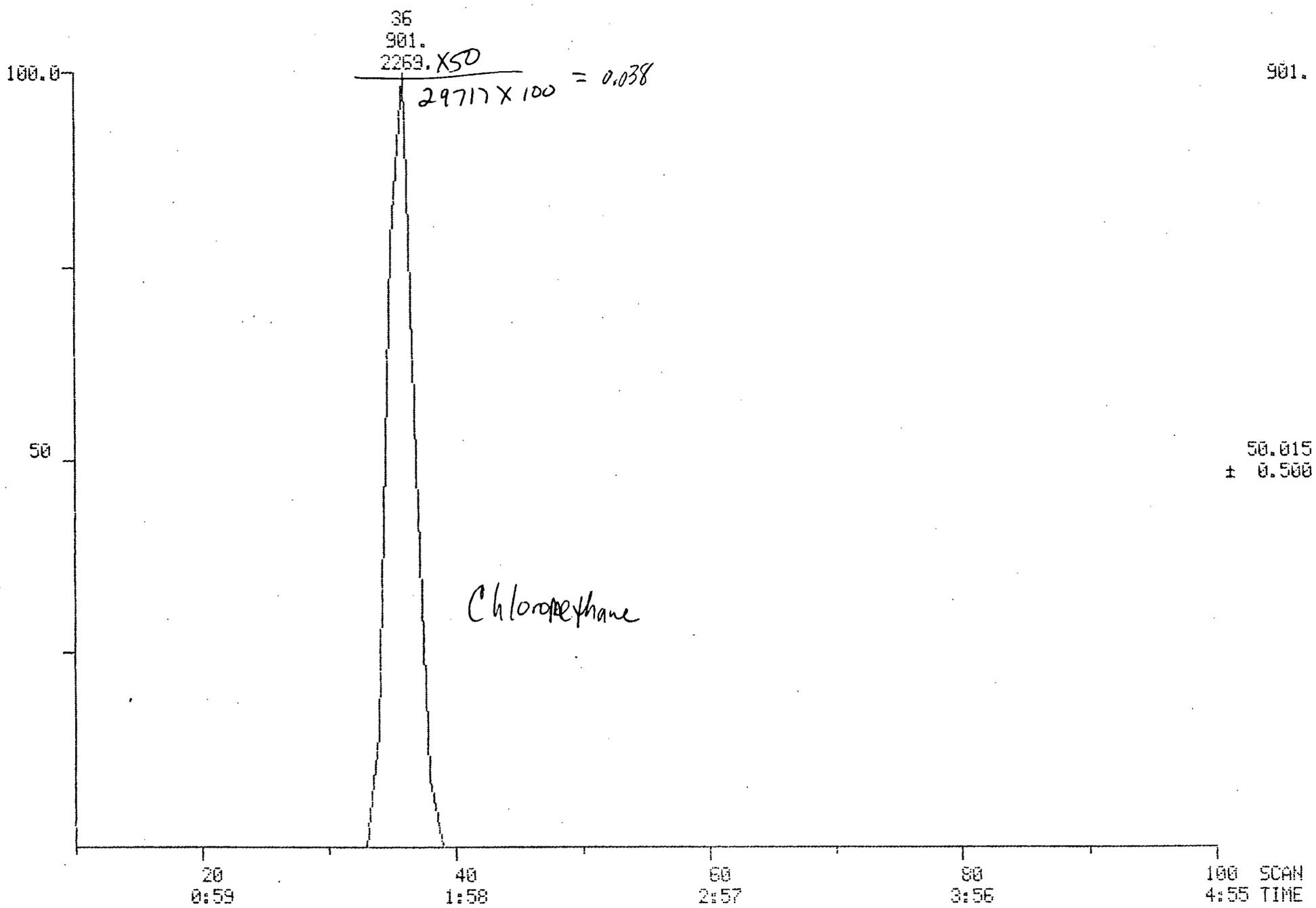
100006 STD (487) 00096

MASS CHROMATOGRAM
05/05/84 20:09:00
SAMPLE:

DATA: CLPSTD13

SCANS 10 TO 100

16000
00097
901.
100ppb STD (587)



UNKNOWN SAMPLE QUANTITATION

OWA REVERSE SEARCH STATUS REPORT

THE INTERNAL STANDARD AREA HAS A DIFFERENCE OF: 520 % OF THE LAST STANDARD RUN COMPOUNDS QUANTITATED BY INTERNAL STANDARD USING PARAMETERS IN LIBRARYUX.

REPORT ENTRY NO.	EXPECTED SCAN	BEST SCAN	FIT	PURITY	LIBRARY ENTRY NO.	PEAKS FOUND	PEAKS QUANT	SATURATED PEAKS
1	147	148	975	925	1	1	1	0
3	51	51	981	816	3	1	1	0
4	61	61	984	643	4	1	1	0
5	74	74	983	769	5	1	1	0
6	100	101	993	956	6	1	1	0
7	123	124	985	942	7	1	1	0
8	140	141	992	918	8	1	1	0
9	162	163	995	898	9	1	1	0
10	174	174	982	936	10	1	1	0
11	184	184	984	921	11	1	1	0
12	194	195	996	399	12	1	1	0
13	196	197	971	813	13	1	1	0
14	217	219	965	922	14	1	1	0
						**		
15	223	225	983	967	15	2	1	0
16	229	231	978	920	16	1	1	0
17	194	195	932	152	17	1	1	0
18	107	108	968	595	18	1	1	0

THE INTERNAL STANDARD AREA HAS A DIFFERENCE OF: 6486 % OF THE LAST STANDARD RUN COMPOUNDS QUANTITATED BY INTERNAL STANDARD USING PARAMETERS IN LIBRARYUY.

REPORT ENTRY NO.	EXPECTED SCAN	BEST SCAN	FIT	PURITY	LIBRARY ENTRY NO.	PEAKS FOUND	PEAKS QUANT	SATURATED PEAKS
						**		
21	294	295	989	465	1	2	1	0
22	250	252	973	910	2	1	1	0
23	253	255	951	804	3	1	1	0
24	260	262	982	886	4	1	1	0
25	269	271	985	626	5	1	1	0
26	273	274	987	433	6	1	1	0
27	273	274	995	218	7	1	1	0
28	271	273	998	257	8	1	1	0

THE INTERNAL STANDARD AREA HAS A DIFFERENCE OF: -69 % OF THE LAST STANDARD RUN COMPOUNDS QUANTITATED BY INTERNAL STANDARD USING PARAMETERS IN LIBRARYUZ.

REPORT ENTRY NO.	EXPECTED SCAN	BEST SCAN	FIT	PURITY	LIBRARY ENTRY NO.	PEAKS FOUND	PEAKS QUANT	SATURATED PEAKS
29	352	355	988	541	1	1	1	0
30	320	322	992	916	2	1	1	0
31	313	315	938	849	3	1	1	0
32	320	322	787	535	4	1	1	0
33	346	349	996	664	5	1	1	0
34	346	349	999	131	6	1	1	0
35	365	368	963	831	7	1	1	0
36	368	372	982	880	8	1	1	0
						**		
37	390	395	998	135	9	2	1	0
38	436	443	984	889	10	1	1	0
39	496	506	986	885	11	1	1	0
40	532	544	975	872	12	1	1	0

1009985TD
(787)

00099

COMPOUND PASSED REVERSE SEARCH BUT WAS BELOW MINIMUM AREA FOR ION QUANTIT

VOA GC/MS SHIFT QUALITY CONTROL REPORT

INSTRUMENT: OWA DATE: 5/5/84 SHIFT: PM OPERATOR: CR

A.L.S. Position	File name	Water or Soil	IS-1 Area Bromochloromethane	SS-1 % D-4 1,2-Dichloroethane	IS-2 Area B-6 Benzene 1-Bromo-2-Chloro Propane	IS-3 Area 1,4-Dichlorobutane	SS-2 % D-8 Toluene	SS-3 % 4-Bromo-fluorobenzene	STATUS
10	²⁷⁵² CLPUDA 29	W	32366	79	49904	51853	92	90	
1	²⁷⁵³ 30		33022	81	52572	53305	91	86	
2	²⁷⁵⁴ 31		32839	77	51117	52778	91	87	78hrs from std.
1	CLPSTD 13		29717	100	45246	31984	100	100	
2	^{BLK} CLPUDA 32		30086	78	45864	33278	109	125	
3	²⁷⁵⁵ 33		31639	88	51163	48435	80	93	
4	²⁷⁵⁶ 34		28267	85	46072	43147	83	87	
5	²⁷⁵⁷ 35		33001	93	53958	53818	76	86	
6	²⁷⁵⁴ 36		30641	85	50534	49832	78	84	

00100

VOA GC/MS SHIFT QUALITY CONTROL REPORT

INSTRUMENT: OWA DATE: 5/5/84 SHIFT: Am OPERATOR: [Signature]

A.L.S. Position	File name	Water or Soil	IS-1 Area Bromochloromethane	SS-1 % D-4 1,2-Dichloroethane	IS-2 Area B-6 Benzene 1-Bromo-2-Chloro Propane	IS-3 Area 1,4-Dichlorobutane	SS-2 % D-8 Toluene	SS-3 % 4-Bromo-fluorobenzene	STATUS
10	CLPSTD 10	W	43990	100	63937	63036	100	100	
1	↓ 11		47635	100	72889	69876	100	100	
2	↓ 12		34782	100	61262	58240	100	100	
3	CLPUDA ^{BLK} 22		6076	50	9435	12083	90	137	Leak
4	2741 23		39589	82	62414	60354	97	96	
5	2741MS 24		32758	82	51961	53883	91	89	
6	2741MSD 25		32529	74	51393	47569	105	103	
7	2741MS ₃ 26		33951	78	52939	54299	94	91	
8	A2750 27		36900	84	57636	59016	93	90	
9	2751 28	↓	36916	80	58406	58277	94	93	

00101

Case No. 2660 Initial Calibration Data - Volatile HSL Compounds
2260 Contractor Cambridge Analytical Contract No. 68-016791
 Instrument Identifier OWA Calibration Date 3/5/84
 Minimum RF for SPCC is 0.050 0.3

COMPOUND	RRT	RF ₂₀	RF ₁₀₀	RF ₂₀₀	RF	SPCC
chloromethane	0.241	0.038	0.062	0.097	0.066	*
bromomethane	0.343	0.211	0.413	0.626	0.417	
dichlorodifluoromethane						
vinyl chloride	0.411	0.076	0.209	0.333	0.206	
chloroethane	0.499	0.091	0.209	0.315	0.205	
methylene chloride	0.679	2.531	2.994	4.259	3.261	
trichlorofluoromethane						
ethene, 1,1-dichloro	0.952	2.377	0.150	1.086	0.738	
ethane, 1,1-dichloro-	1.102	0.985	1.743	2.478	1.735	*
1,2-trans-dichloroethene	1.180	0.396	0.863	1.289	0.849	
chloroform	1.250	1.772	2.960	4.259	2.997	
ethane, 1,2-dichloro-	1.332	1.131	2.083	2.999	2.071	
ethane, 1,1,1-trichloro-	1.477	1.469	2.691	4.062	2.741	
carbon tetrachloride	1.548	1.205	2.542	4.343	2.697	
bromodichloromethane	1.559	0.603	1.892	3.116	1.870	
propane, 1,2-dichloro-	0.851	0.350	0.657	0.865	0.624	
1,3-trans-dichloropropene	0.862	0.407	0.899	1.205	0.837	
trichloroethylene	0.889	0.738	1.096	1.332	1.055	
chlorodibromomethane	0.924	0.132	0.667	1.125	0.641	
benzene	0.917	1.011	1.737	2.335	1.694	
ethane, 1,1,2-trichloro-	0.931	0.451	0.707	0.957	0.705	
1,3-cis-dichloropropene	0.928	0.312	0.765	1.194	0.757	*
2-chloroethyl vinyl ether						*
bromoform	0.838	0.034	0.351	0.775	0.387	*
ethane, 1,1,2,2-tetrachloro-	0.983	0.708	1.071	1.430	1.076	*
ethene, tetrachloro-	0.983	0.806	1.244	1.541	1.197	
toluene	1.046	1.383	1.523	1.882	1.596	
chlorobenzene	1.110	1.830	2.483	3.288	2.534	*
ethylbenzene	1.241	0.485	1.149	1.546	1.060	
acetone	0.727	0.346	0.197	0.327	0.290	
2-butanone	1.318	0.150	0.270	0.466	0.295	
carbonyl sulfide	0.834	1.080	1.688	2.684	1.815	
2-hexanone	0.907	0.330	0.516	0.744	0.530	
4-methyl-2-pentanone	0.907	0.330	0.516	0.744	0.530	
styrene	1.517	0.939	2.045	2.773	1.919	
vinyl acetate						
o-xylene	1.606	0.552	1.368	1.843	1.254	

RRT - Average Relative Retention Time
 RF - Response Factor (subscript is the amount of nanograms)
 RF - Average Response Factor
 SPCC - System Performance Check Compounds (those compounds flagged with an *)



Cambridge Analytical Associates

RAW QC PACKET

00104

GC/MS PERFORMANCE STANDARD

Bromofluorobenzene

CASE NO. 2660 CONTRACTOR Cambridge Anal. CONTRACT NO. 68-01-6791
 INSTRUMENT ID OWA DATE 5/5/84 TIME 10:42
 RUN NUMBER BFBDI QC REPORT NO. 011 ANALYST EAC

TUNE CHECK:

m/e	Ion Abundance Criteria	% Relative Abundance
50	15 - 40% of the base peak	19
75	30 - 60% of the base peak	60
95	Base peak, 100% relative abundance	100
96	5 - 9% of the base peak	5
173	Less than 1% of the base peak	—
174	Greater than 50% of the base peak	72
175	5 - 9% of mass 174	4 (6) ¹
176	Greater than 95%, but less than 101% of 174	70 (99) ¹
177	5 - 9% of mass 176	3 (5) ²

¹Value in parenthesis is % of mass 174.

²Value in parenthesis is % of mass 176.

Comments:

FORM VI

Revision Date 1/83

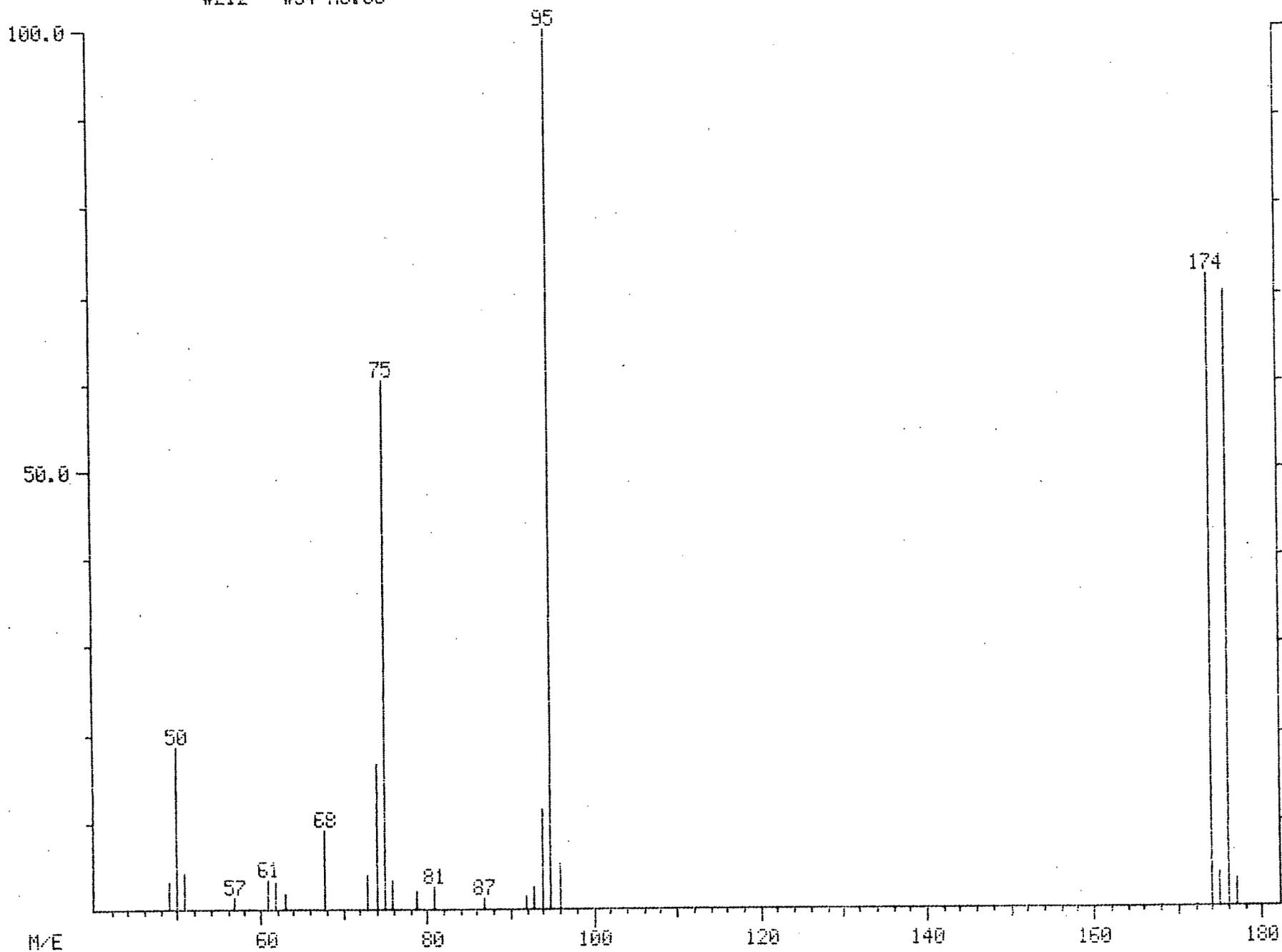
00105

MASS SPECTRUM
05/05/84 10:42:00 + 10:25
SAMPLE:
#212 - #54 X3.05

DATA: BFB01 #212

BASE M/E: 95
RIC: 58848.

00106



14288.
10.

MASS LIST
05/05/84 10:42:00 + 10:25

DATA: EFRDI # 212

BASE M/E: 95
RIC: 58048.

SAMPLE:

#212 - #54 X3.05

MASS	% RA	MIN INTEN:	0.	MAX INTEN:	14288.
49	0.00	MINIMA			
177	# 0	MAXIMA			
49	3.18				
50	18.62				
51	4.26				
57	1.29				
61	3.30				
62	3.14				
63	1.85				
68	9.13				
73	3.93				
74	16.77				
75	60.47				
76	3.48				
79	2.07				
81	2.51				
87	1.29				
92	1.47				
93	2.72				
94	11.55				
95	100.00				
96	5.21				
174	71.89				
175	4.02				
176	70.21				
177	3.18				

00107

GC/MS PERFORMANCE STANDARD

Bromofluorobenzene

CASE NO. 2660 CONTRACTOR Cambridge Anal. CONTRACT NO. 68-01-6791
 INSTRUMENT ID OWA DATE 5/5/84 TIME 19:46
 RUN NUMBER BFB DI 1 QC REPORT NO. 011 ANALYST EC

TUNE CHECK:

<u>m/e</u>	<u>Ion Abundance Criteria</u>	<u>% Relative Abundance</u>
50	15 - 40% of the base peak	20
75	30 - 60% of the base peak	60
95	Base peak, 100% relative abundance	100
96	5 - 9% of the base peak	5
173	Less than 1% of the base peak	—
174	Greater than 50% of the base peak	86
175	5 - 9% of mass 174	5 (5) ¹
176	Greater than 95%, but less than 101% of 174	86 (100) ²
177	5 - 9% of mass 176	4 (5) ²

¹Value in parenthesis is % of mass 174.
²Value in parenthesis is % of mass 176.

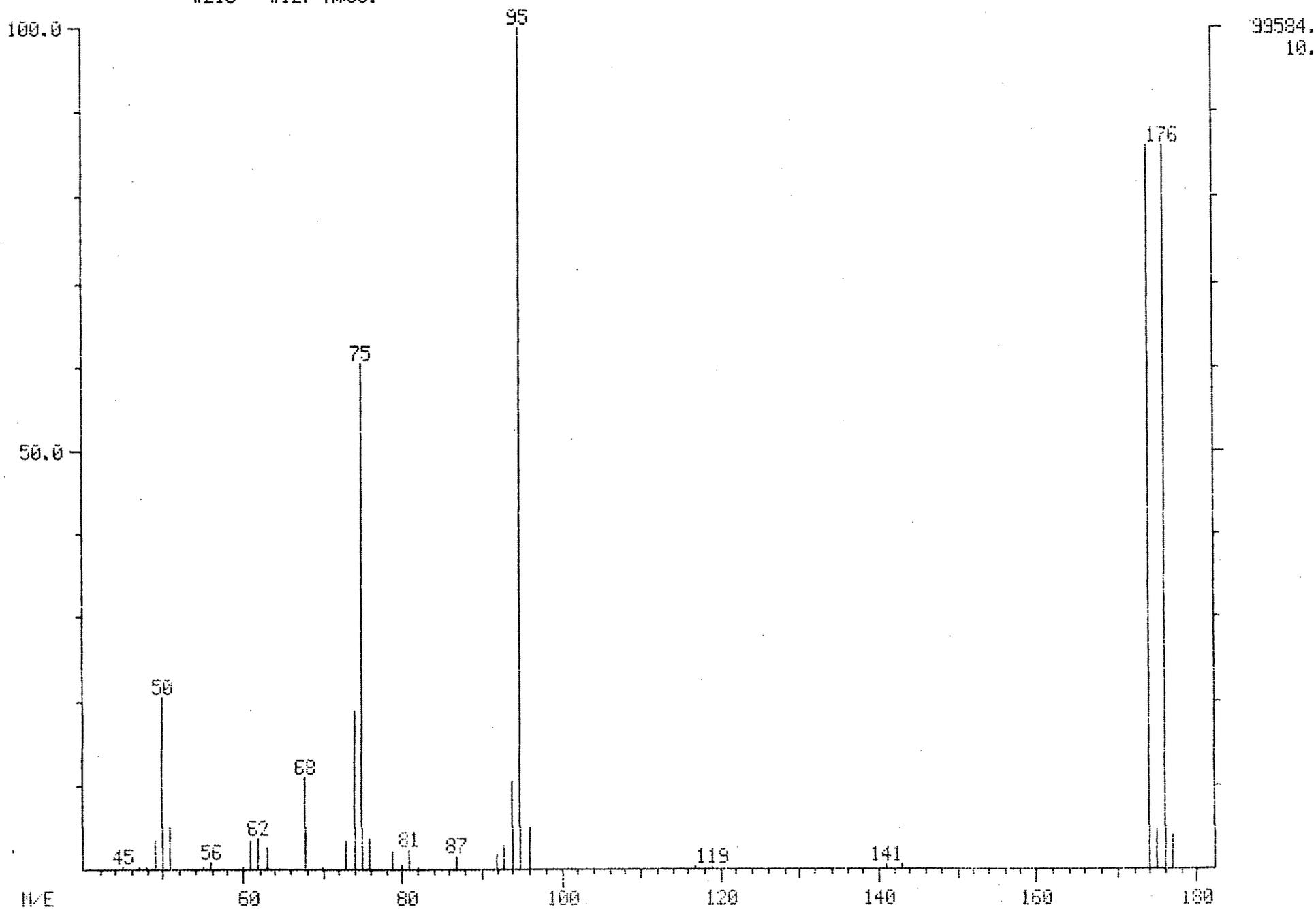
Comments:

MASS SPECTRUM
05/05/84 19:46:00 + 10:19
SAMPLE:
#210 - #127 X*35.

DATA: BFBDI1 #210

BASE M/E: 95
RIC: 445440.

00109



MASS LIST
05/05/84 19:46:00 + 10:19
SAMPLE:

DATA: BFED11 # 210

BASE M/E: 95
RIC: 445440.

#210 - #127 X*35.

MASS	% RA	MIN INTEN:	MAX INTEN:
41	0.00	0.	99584.
177 #	0		
41	0.00		
45	0.38		
47	0.27		
48	0.19		
49	3.31		
50	20.31		
51	4.99		
55	0.15		
56	0.88		
60	0.33		
61	3.45		
62	3.68		
63	2.60		
68	10.89		
70	0.35		
72	0.10		
73	3.43		
74	18.77		
75	60.41		
76	3.69		
79	2.19		
80	0.51		
81	2.30		
82	0.38		
87	1.53		
92	1.88		
93	2.90		
94	10.54		
95	100.00		
96	4.96		
117	0.21		
119	0.37		
141	0.46		
143	0.41		
174	85.86		
175	4.60		
176	85.99		
177	3.94		

RIC
05/05/84 21:01:00
SAMPLE:

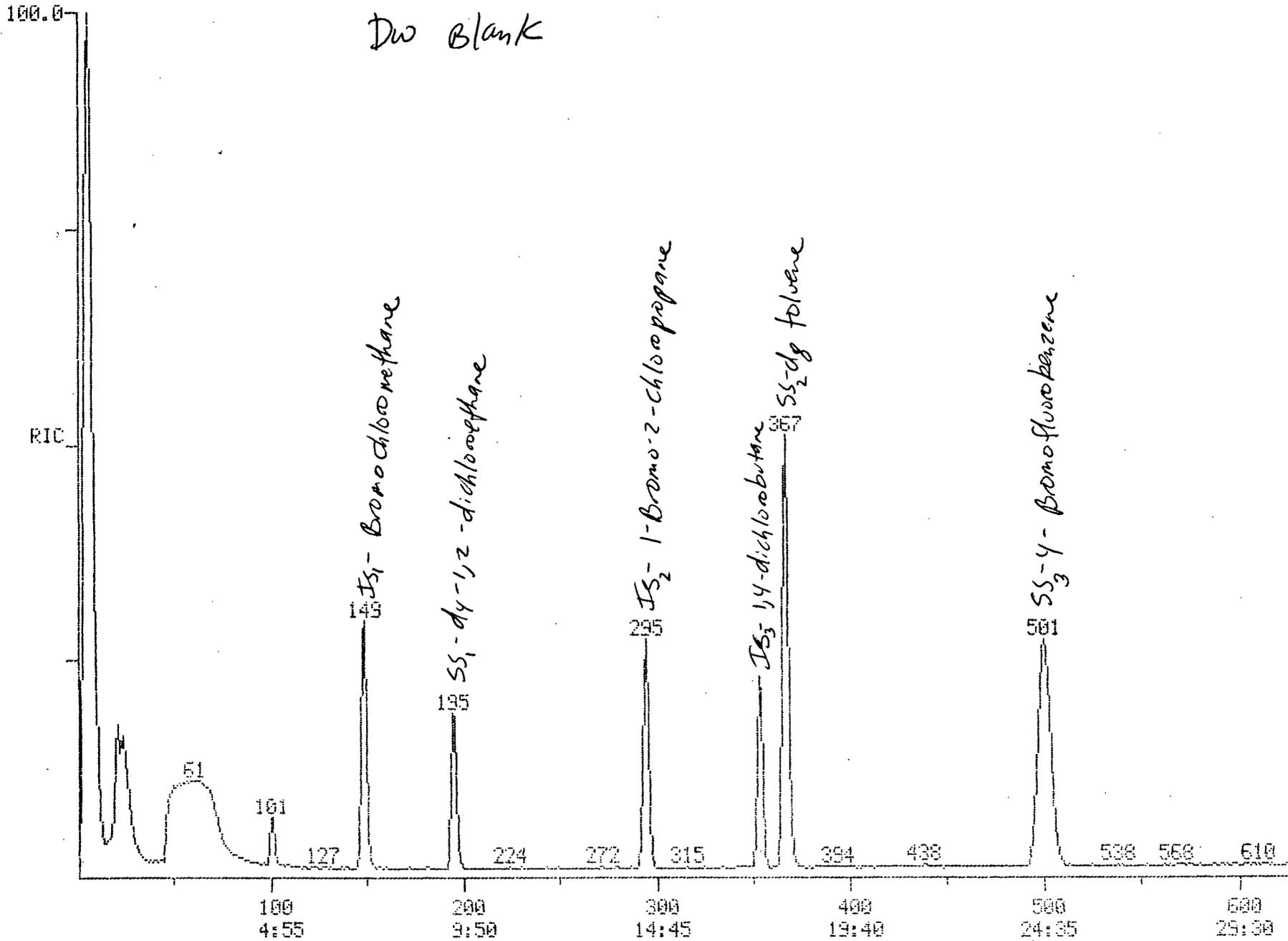
DATA: CLPVOA32

SCANS 1 TO 625

CASE 2060
Dw Blank

128128.

00111



VOA blank (104)

SCAN
TIME

FINNIGAN ORGANICS IN WATER ANALYZER
QUANTITATION REPORT FILE: CLPVOA32

DATA: CLPVOA32.TI
05/05/84 21:01:00

SAMPLE:

SUBMITTED BY:

ANALYST:

AMOUNT=AREA(HGHT) * REF. AMNT / (REF. AREA(HGHT) * RESP. FACT)
RESP. FAC. FROM LIBRARY ENTRY

- NO NAME
- 1 BROMOCHLOROMETHANE (INTERNAL STANDARD)
 - 2 CHLOROMETHANE
 - 3 BROMOMETHANE
 - 4 VINYL CHLORIDE
 - 5 CHLOROETHANE
 - 6 METHYLENE CHLORIDE
 - 7 CARBON DISULFIDE
 - 8 1,1 DICHLOROETHYLENE
 - 9 1,1 DICHLOROETHANE
 - 10 TRANS 1,2 DICHLOROETHYLENE
 - 11 CHLOROFORM
 - 12 D4-1,2 DICHLOROETHANE (SURROGATE STANDARD)
 - 13 1,2 DICHLOROETHANE
 - 14 1,1,1 TRICHLOROETHANE
 - 15 CARBON TETRACHLORIDE
 - 16 BROMODICHLOROMETHANE
 - 17 2-BUTANONE (MEK)
 - 18 ACETONE
 - 19 ACRYLONITRILE
 - 20 ACROLEIN
 - 21 1-BROMO-2-CHLOROPROPANE (INTERNAL STANDARD)
 - 22 1,2 DICHLOROPROPANE
 - 23 TRANS 1,3-DICHLOROPROPENE
 - 24 TRICHLOROETHYLENE
 - 25 BENZENE
 - 26 1,1,2-TRICHLOROETHANE
 - 27 CIS 1,3-DICHLOROPROPENE
 - 28 DIBROMOCHLOROMETHANE
 - 29 1,4 DICHLOROBUTANE (INTERNAL STANDARD)
 - 30 4-METHYL 2-PENTANONE (MIBK)
 - 31 BROMOFORM
 - 32 2-HEXANONE (MPK)
 - 33 TETRACHLOROETHYLENE
 - 34 1,1,2,2 TETRACHLOROETHANE
 - 35 D8-TOLUENE (SURROGATE STANDARD)
 - 36 TOLUENE
 - 37 CHLOROBENZENE
 - 38 ETHYLBENZENE
 - 39 4-BROMOFLUOROBENZENE (SURROGATE STANDARD)
 - 40 STYRENE
 - 41 O-XYLENE

NO	M/E	SCAN	TIME	REF	RRT	METH	AREA(HGHT)	AMOUNT	%TOT
1	130	148	7:17	1	1.000	A BB	30086.	50.000 UG/L	10.70
2	NOT FOUND								
3	NOT FOUND								

NOA blank (294)

00112

NO	M/E	SCAN	TIME	REF	RRT	METH	AREA(HGHT)	AMOUNT	%TOT
4		NOT FOUND							
5		NOT FOUND							
6	84	101	4:58	1	0.682	A BB	5272.	2.374 UG/L	0.51 <i>BLK</i>
7		NOT FOUND							
8		NOT FOUND							
9		NOT FOUND							
10		NOT FOUND							
11		NOT FOUND							
12	102	195	9:35	1	1.318	A BB	3475.	78.009 PRCNT	16.69 ✓
13		NOT FOUND							
14		NOT FOUND							
15		NOT FOUND							
16		NOT FOUND							
17		NOT FOUND							
18	43	108	5:19	1	0.730	A BB	382.	3.114 UG/L	0.67 <i>BLK</i>
19		NOT FOUND							
20		NOT FOUND							
21	77	295	14:30	21	1.000	A BB	45864.	50.000 UG/L	10.70
22		NOT FOUND							
23		NOT FOUND							
24		NOT FOUND							
25		NOT FOUND							
26		NOT FOUND							
27		NOT FOUND							
28		NOT FOUND							
29	55	353	17:21	29	1.000	A BB	33278.	50.000 UG/L	10.70
30		NOT FOUND							
31		NOT FOUND							
32		NOT FOUND							
33		NOT FOUND							
34		NOT FOUND							
35	100	367	18:03	29	1.040	A BB	70082.	109.001 PRCNT	23.32 ✓
36		NOT FOUND							
37		NOT FOUND							
38		NOT FOUND							
39	95	501	24:38	29	1.419	A BB	68245.	124.832 PRCNT	26.71 ✓
40		NOT FOUND							
41		NOT FOUND							

VOAblank (3084)

00113

UNKNOWN SAMPLE QUANTITATION

OWA REVERSE SEARCH STATUS REPORT

THE INTERNAL STANDARD AREA HAS A DIFFERENCE OF: 528 % OF THE LAST STANDARD RUN
COMPOUNDS QUANTITATED BY INTERNAL STANDARD USING PARAMETERS IN LIBRARYUX

REPORT ENTRY NO.	EXPECTED SCAN	BEST SCAN	FIT	PURITY	LIBRARY ENTRY NO.	PEAKS FOUND	PEAKS QUANT	SATURATED PEAKS
1	148	149	978	916	1	1	1	0
6	101	101	956	742	6	1	1	0
12	195	195	1000	796	12	1	1	0
18	108	108	762	73	18	1	1	0

THE INTERNAL STANDARD AREA HAS A DIFFERENCE OF: 6575 % OF THE LAST STANDARD RUN
COMPOUNDS QUANTITATED BY INTERNAL STANDARD USING PARAMETERS IN LIBRARYUY

REPORT ENTRY NO.	EXPECTED SCAN	BEST SCAN	FIT	PURITY	LIBRARY ENTRY NO.	PEAKS FOUND	PEAKS QUANT	SATURATED PEAKS
21	295	295	974	500	1	1	1	0

THE INTERNAL STANDARD AREA HAS A DIFFERENCE OF: -73 % OF THE LAST STANDARD RUN
COMPOUNDS QUANTITATED BY INTERNAL STANDARD USING PARAMETERS IN LIBRARYUZ

REPORT ENTRY NO.	EXPECTED SCAN	BEST SCAN	FIT	PURITY	LIBRARY ENTRY NO.	PEAKS FOUND	PEAKS QUANT	SATURATED PEAKS
29	355	353	987	520	1	1	1	0
35	368	367	967	863	7	1	1	0
39	506	501	984	861	11	1	1	0

NUMBER OF COMPOUNDS IDENTIFIED 8

DATA PROCESSING OF CLPV0A32 COMPLETED ON 5/05/84 21:32:16

VOA blank (4084) 00114

RIC
05/05/84 12:52:00
SAMPLE:

DATA: CLPVOA24

SCANS 1 TO 650

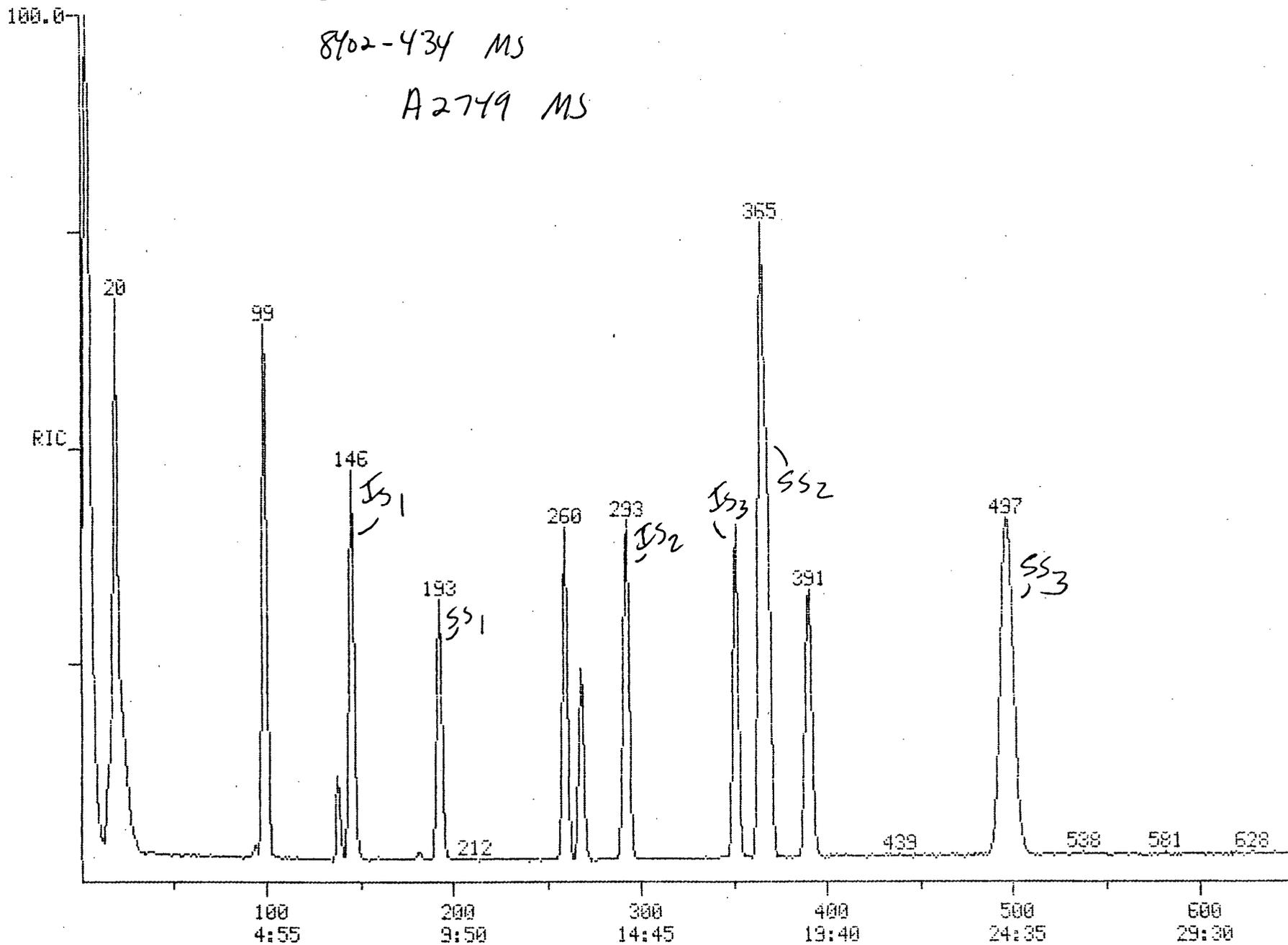
00115

CASE 2660

8902-434 MS

A2749 MS

102528.



A2749 MS (1984)

SCAN
TIME

FINNIGAN ORGANICS IN WATER ANALYZER
 QUANTITATION REPORT FILE: CLPVOA24

DATA: CLPVOA24.TI
 05/05/84 12:52:00
 SAMPLE:
 SUBMITTED BY:

ANALYST:

AMOUNT=AREA(HGHT) * REF. AMNT / (REF. AREA(HGHT) * RESP. FACT)
 RESP. FAC. FROM LIBRARY ENTRY

- | NO | NAME |
|----|---|
| 1 | BROMOCHLOROMETHANE (INTERNAL STANDARD) |
| 2 | CHLOROMETHANE |
| 3 | BROMOMETHANE |
| 4 | VINYL CHLORIDE |
| 5 | CHLOROETHANE |
| 6 | METHYLENE CHLORIDE |
| 7 | CARBON DISULFIDE |
| 8 | 1,1 DICHLOROETHYLENE |
| 9 | 1,1 DICHLOROETHANE |
| 10 | TRANS 1,2 DICHLOROETHYLENE |
| 11 | CHLOROFORM |
| 12 | D4-1,2 DICHLOROETHANE (SURROGATE STANDARD) |
| 13 | 1,2 DICHLOROETHANE |
| 14 | 1,1,1 TRICHLOROETHANE |
| 15 | CARBON TETRACHLORIDE |
| 16 | BROMODICHLOROMETHANE |
| 17 | 2-BUTANONE (MEK) |
| 18 | ACETONE |
| 19 | ACRYLONITRILE |
| 20 | ACROLEIN |
| 21 | 1-BROMO-2-CHLOROPROPANE (INTERNAL STANDARD) |
| 22 | 1,2 DICHLOROPROPANE |
| 23 | TRANS 1,3-DICHLOROPROPENE |
| 24 | TRICHLOROETHYLENE |
| 25 | BENZENE |
| 26 | 1,1,2-TRICHLOROETHANE |
| 27 | CIS 1,3-DICHLOROPROPENE |
| 28 | DIBROMOCHLOROMETHANE |
| 29 | 1,4 DICHLOROBUTANE (INTERNAL STANDARD) |
| 30 | 4-METHYL 2-PENTANONE (MIEK) |
| 31 | BROMOFORM |
| 32 | 2-HEXANONE (MPK) |
| 33 | TETRACHLOROETHYLENE |
| 34 | 1,1,2,2 TETRACHLOROETHANE |
| 35 | D8-TOLUENE (SURROGATE STANDARD) |
| 36 | TOLUENE |
| 37 | CHLOROBENZENE |
| 38 | ETHYLBENZENE |
| 39 | 4-BROMOFLUOROBENZENE (SURROGATE STANDARD) |
| 40 | STYRENE |
| 41 | O-XYLENE |

NO	M/E	SCAN	TIME	REF	RRT	METH	AREA(HGHT)	AMOUNT	%TOT
1	130	146	7:11	1	1.000	A BB	32758.	50.000 UG/L	9.08
2	NOT FOUND								
3	NOT FOUND								

A2789MS(284)

00116

NO	M/E	SCAN	TIME	REF	RRT	METH	AREA(HGHT)	AMOUNT	%TOT
4		NOT FOUND							
5		NOT FOUND							
6	84	99	4:52	1	0.678	A BB	44971.	22.927 UG/L	4.16 <i>BY</i>
7		NOT FOUND							
8	96	139	6:50	1	0.952	A BB	5847.	11.897 UG/L	2.16
9		NOT FOUND							
10		NOT FOUND							
11		NOT FOUND							
12	102	193	9:29	1	1.322	A BB	4472.	82.305 PRCNT	14.95 ✓
13		NOT FOUND							
14		NOT FOUND							
15		NOT FOUND							
16		NOT FOUND							
17		NOT FOUND							
18	43	106	5:13	1	0.726	A BB	782.	6.061 UG/L	1.10 <i>BY</i>
19		NOT FOUND							
20		NOT FOUND							
21	77	293	14:24	21	1.000	A BB	51961.	50.000 UG/L	9.08
22		NOT FOUND							
23		NOT FOUND							
24	130	260	12:47	21	0.887	A BB	24930.	21.886 UG/L	3.98
25	78	268	13:11	21	0.915	A BB	45119.	24.992 UG/L	4.54
26		NOT FOUND							
27		NOT FOUND							
28		NOT FOUND							
29	55	352	17:18	29	1.000	A BB	53883.	50.000 UG/L	9.08
30		NOT FOUND							
31		NOT FOUND							
32		NOT FOUND							
33		NOT FOUND							
34		NOT FOUND							
35	100	365	17:57	29	1.037	A BB	77621.	91.086 PRCNT	16.54 ✓
36	92	368	18:06	29	1.045	A BB	45118.	27.494 UG/L	4.99
37	112	391	19:13	29	1.111	A BB	60753.	22.704 UG/L	4.12
38		NOT FOUND							
39	95	498	24:29	29	1.415	A BB	78557.	89.242 PRCNT	16.21 ✓
40		NOT FOUND							
41		NOT FOUND							

A 2749 MS(384)

00117

THE INTERNAL STANDARD AREA HAS A DIFFERENCE OF: 584 % OF THE LAST STANDARD RUN
COMPOUNDS QUANTITATED BY INTERNAL STANDARD USING PARAMETERS IN LIBRARYUX

REPORT ENTRY NO.	EXPECTED SCAN	BEST SCAN	FIT	PURITY	LIBRARY ENTRY NO.	PEAKS FOUND	PEAKS QUANT	SATURATED PEAKS
1	147	146	983	908	1	1	1	0
6	100	99	992	935	6	1	1	0
8	140	139	983	754	8	1	1	0
12	194	193	996	754	12	1	1	0
18	107	106	761	83	18	1	1	0

THE INTERNAL STANDARD AREA HAS A DIFFERENCE OF: 7463 % OF THE LAST STANDARD RUN
COMPOUNDS QUANTITATED BY INTERNAL STANDARD USING PARAMETERS IN LIBRARYUY

REPORT ENTRY NO.	EXPECTED SCAN	BEST SCAN	FIT	PURITY	LIBRARY ENTRY NO.	PEAKS FOUND	PEAKS QUANT	SATURATED PEAKS
21	294	293	985	446 -	1	2	1	0
24	260	260	979	864	4	1	1	0
25	269	268	973	837	5	1	1	0

THE INTERNAL STANDARD AREA HAS A DIFFERENCE OF: -55 % OF THE LAST STANDARD RUN
COMPOUNDS QUANTITATED BY INTERNAL STANDARD USING PARAMETERS IN LIBRARYUZ

REPORT ENTRY NO.	EXPECTED SCAN	BEST SCAN	FIT	PURITY	LIBRARY ENTRY NO.	PEAKS FOUND	PEAKS QUANT	SATURATED PEAKS
29	352	352	982	509	1	1	1	0
35	365	365	972	822	7	1	1	0
36	368	368	972	764	8	1	1	0
37	390	391	998	130 -	9	2	1	0
39	496	497	988	831	11	1	1	0

NUMBER OF COMPOUNDS IDENTIFIED 13

DATA PROCESSING OF CLPVOA24 COMPLETED ON 5/05/84 14:00:35

A2749MS (40/4)

00118

RIC
05/05/84 13:39:00
SAMPLE:

DATA: CLPVOA25

SCANS 1 TO 650

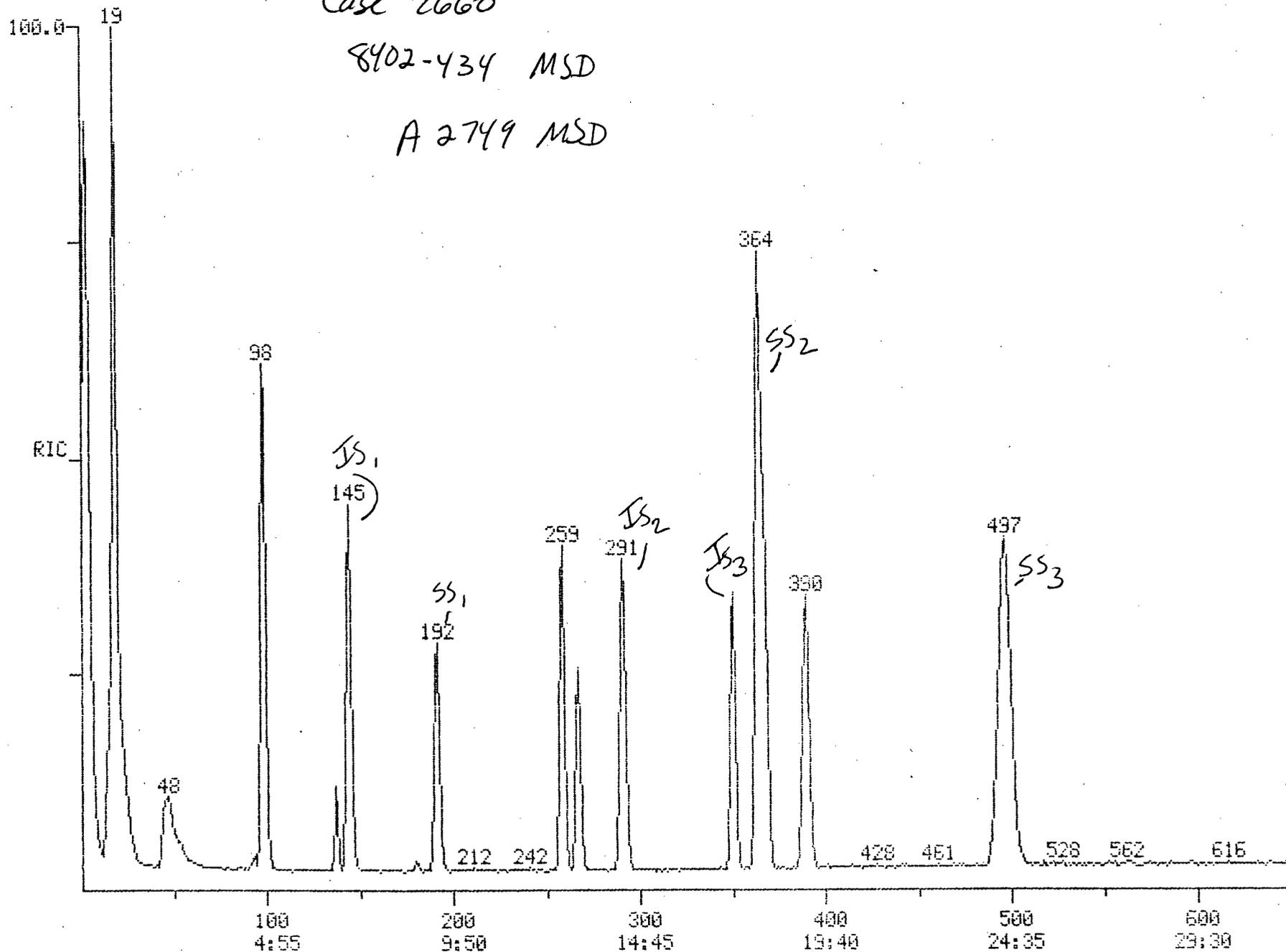
00119

Case 2660

8402-434 MSD

A 2749 MSD

107264.



A 2749 MSD (1074)

SCAN
TIME

100.0-
19

RIC
05/05/84 13:39:00
SAMPLE:

Case 2660

8402-434 MSD

A 2749 MSD

DATA: CLPUDR25

SCANS 1 TO 650

107264.

00119

REF	RRT	METH	AREA(HGHT)	AMOUNT	%TOT
1	0.676	A BB	45465.	23.342 UG/L	4.07 <i>BLK</i>
1	0.952	A BB	6418.	13.151 UG/L	2.29
1	1.324	A BB	4004.	74.211 PRCNT	12.95 ✓
E1	1.000	A BB	51393.	50.000 UG/L	8.72
E1	0.890	A BB	25234.	22.398 UG/L	3.91
E1	0.918	A BB	47042.	26.345 UG/L	4.60
E9	1.000	A BB	47569.	50.000 UG/L	8.72
E9	1.037	A BB	79058.	105.087 PRCNT	18.33 ✓
E9	1.046	A BB	41985.	28.981 UG/L	5.06
E9	1.111	A BB	62725.	26.553 UG/L	4.63
E9	1.416	A BB	80158.	103.147 PRCNT	17.99 ✓

A2749MSD (384)

00121

THE INTERNAL STANDARD AREA HAS A DIFFERENCE OF: 579 % OF THE LAST STANDARD RUN
COMPOUNDS QUANTITATED BY INTERNAL STANDARD USING PARAMETERS IN LIBRARYUX

REPORT ENTRY NO.	EXPECTED SCAN	BEST SCAN	FIT	PURITY	LIBRARY ENTRY NO.	PEAKS FOUND	PEAKS QUANT	SATURATED PEAKS
1	147	145	986	908	1	1	1	0
6	100	98	995	944	6	1	1	0
8	140	138	981	763	8	1	1	0
12	194	192	996	752	12	1	1	0
COMPOUND PASSED REVERSE SEARCH BUT WAS BELOW MINIMUM AREA FOR ION QUANTIT								
18	107	105	755	29	18	1	0	0

THE INTERNAL STANDARD AREA HAS A DIFFERENCE OF: 7380 % OF THE LAST STANDARD RUN
COMPOUNDS QUANTITATED BY INTERNAL STANDARD USING PARAMETERS IN LIBRARYUY

REPORT ENTRY NO.	EXPECTED SCAN	BEST SCAN	FIT	PURITY	LIBRARY ENTRY NO.	PEAKS FOUND	PEAKS QUANT	SATURATED PEAKS
21	294	291	984	474	1	2	1	0
24	260	259	983	873	4	1	1	0
25	269	267	975	853	5	1	1	0

THE INTERNAL STANDARD AREA HAS A DIFFERENCE OF: -61 % OF THE LAST STANDARD RUN
COMPOUNDS QUANTITATED BY INTERNAL STANDARD USING PARAMETERS IN LIBRARYUZ

REPORT ENTRY NO.	EXPECTED SCAN	BEST SCAN	FIT	PURITY	LIBRARY ENTRY NO.	PEAKS FOUND	PEAKS QUANT	SATURATED PEAKS
29	352	351	985	502	1	1	1	0
35	365	364	970	824	7	1	1	0
36	368	367	975	756	8	1	1	0
37	390	390	1000	129 -	9	2	1	0
39	496	497	991	828	11	1	1	0

NUMBER OF COMPOUNDS IDENTIFIED 13
DATA PROCESSING OF CLPVOA25 COMPLETED ON 5/05/84 14:47:41

A2749MSD (484)

00122

REGIONAL REVIEW OF UNCONTROLLED HAZARDOUS WASTE SITE CONTRACT
LABORATORY DATA PACKAGE

TO: U.S. Environmental Protection Agency
Sample Management Office (SMO)
P.O. Box 818
Alexandria, Virginia 22313

The hardcopied (laboratory name) Cambridge Analytical Assoc. data package received at Region I has been reviewed and the quality assurance and performance data summarized. The data reviewed included:

Case No.	SMD Sample No.	SMD Sample No.	SMD Sample No.
<u>2660</u>	<u>A2749</u>	<u>A2750</u>	<u>A2751</u>
_____	<u>A2752</u>	<u>A2753</u>	<u>A2754</u>
_____	<u>A2755</u>	<u>A2756</u>	<u>A2757</u>
_____	_____	_____	_____

Contract No. 68-01-6791 requires that specific analytical work be done and that associated reports be provided by the contractor to the Regions, EMSL-LV, and SMO. The general criteria used to determine the performance were based on an examination of:

- Data completeness
- Spectra matching quality
- Surrogate spike results
- Matrix spike results
- Duplicate analysis results
- Blank analysis results
- DFTPP and BFB performance results

The data review forms for each of the above review items are contained within the body of this memo.

Comments: Chloroform, Methylene Chloride, Toluene and Acetone
should be rejected due to blank contamination. Also, although
some surrogate spikes exceeded the limits for recovery, these
limits are only advisory at this time and should not
justify using this as a basis for rejecting data.

I. DATA COMPLETENESS CHECKLIST

- Organics analysis data sheets
- Base/neutral - sample chromatograms
- Acid - sample chromatograms
- VOA - sample chromatograms
- Pesticide - sample chromatograms
- Sample spectra - priority pollutants and non-priority pollutants
- Library reference spectra
- Blank analysis data
- Duplicate analysis: one duplicate analysis of sample A2749 was reported as required by contract.
- Spike data
- DFTPP criteria forms, spectra and listings
- BFB criteria forms, spectra and listings
- Base/neutral - standard reference spectra and chromatograms
- Acid - standard reference spectra and chromatograms
- VOA - standard reference spectra and chromatograms
- Pesticide - standard chromatogram
- Base/neutral sensitivity test
- Acid sensitivity test
- Tailing factor data

Remarks: Standard reference spectra or library reference
was missing

II. SPECTRA MATCHING QUALITY

_____ The spectra were examined and found to be of good matching quality.

_____ The spectra were examined and found to be of poor matching quality due to:

Specified Hazardous Substances List (HSL) Compounds Detected:

Sample No.	Compound Name	Comments
_____	_____	_____
_____	_____	_____
_____	_____	_____
_____	_____	_____
_____	_____	_____
_____	_____	_____
_____	_____	_____
_____	_____	_____
_____	_____	_____
_____	_____	_____
_____	_____	_____
_____	_____	_____
_____	_____	_____
_____	_____	_____
_____	_____	_____

Tentatively Identified Compounds Detected:

Sample No.	Compound Name	Comments
_____	_____	_____
_____	_____	_____
_____	_____	_____
_____	_____	_____
_____	_____	_____
_____	_____	_____
_____	_____	_____
_____	_____	_____
_____	_____	_____
_____	_____	_____
_____	_____	_____
_____	_____	_____
_____	_____	_____
_____	_____	_____
_____	_____	_____

Remarks: Didn't find other spectra for specific compounds in files
but spectra matches well w/ EPA/NIH Mass Spec Ref Data Base

III. SURROGATE SPIKES

The recoveries of the surrogate spiking compounds were reviewed. The average recovery results, standard deviations, etc., are listed below. Asterisked values may indicate quality control problems.

Fraction	Surrogate Compound	Acceptable Range (%)	Average Recovery (%)	+ Standard Deviation	Minimum Recovery (%)	Maximum Recovery (%)	Number Out of Control	Total Number of Samples
VOA	Benzene-d ₆	70 - 130						
VOA	Toluene-d ₈	70 - 130 84 - 114	87%	7.6	75%	97%	4	9
Acid	Phenol-d ₅	30 - 100						
Acid	2-fluorophenol	30 - 100						
B/N	Nitrobenzene-d ₅	40 - 120						
B/N	2-fluorobiphenyl	40 - 120						
VOA	RFB	63-127*	89%	3.8	84%	96%	0	9
VOA	D _{1,1,2} Dichloro Ethane	90-130*	84%	4.1	79%	93%	8	9

18

Remarks: * Advisory Limit Data suspect due to 44% of samples for Toluene out of control. Also, although the limits are advisory, all but one sample for DCE was out of control. For this compound, even the QA surrogates were out.

IV. MATRIX SPIKE RESULTS

The matrix spike results (MSR) for each parameter group were evaluated. The parameters that were reported are listed below along with the MSR guidelines and amount of spike added. A double asterisk (**) indicates outliers.

Compound	Acceptable Range (%)	Average Recovery (%)	± Standard Deviation	Number Out of Control	Total Number	Outliers
<u>Volatiles</u>						
Chlorobenzene	60 - 150	99%	7.5	0	2	
Toluene	40 - 190	103%	3	0	2	
Benzene	70 - 200	102%	2.5	0	2	
<u>Base/Neutrals</u>						
1,2,4-trichlorobenzene	50 - 200					
Acenaphthene	35 - 200					
2,4-dinitrotoluene	25 - 200					
Di-n-butylphthalate	50 - 180					
Pyrene	50 - 150					
N-nitrosodi-n-propylamine	20 - 100					
1,4-dichlorobenzene	15 - 200					
<u>Acids</u>						
Pentachlorophenol	40 - 140					
Phenol	50 - 200					
2-chlorophenol	40 - 150					
p-chloro-m-cresol	40 - 120					
4-nitrophenol	90 - 200					
<u>Pesticides</u>						
Heptachlor	70 - 150					
Aldrin	80 - 150					
Dieldrin	85 - 150					

**Not within specified criteria.

Remarks:

V. DUPLICATE ANALYSIS RESULTS

The relative percent difference (RPD) for each parameter group was evaluated. The duplicate analysis RPD acceptance criteria should be:

<u>Fraction</u>	<u>Maximum Acceptable Percent Difference</u>
Volatile	15%
Base/Neutral	50%
Acid	40%

The RPDs exceeding the maximum acceptable percent difference were:

<u>Fraction</u>	<u>Compound</u>	<u>Actual RPD</u>	<u>Concentration</u>	
			<u>Sample</u>	<u>Duplicate</u>
Volatile	<u>Chlorobenzene</u>	<u>15.8</u>	<u>22.7</u>	<u>26.6</u>
Base/Neutral	_____	_____	_____	_____
Acid	_____	_____	_____	_____

Each duplicate analysis was examined in reference to compounds detected in each analysis. Those compounds which were not common to each analysis for the duplicate sample are listed below:

<u>Fraction</u>	<u>Sample No.</u>	<u>Compound</u>	<u>Concentration</u>
_____	_____	_____	_____
_____	_____	_____	_____
_____	_____	_____	_____

Remarks: My calculations give me an RPD of 7.9. In either case, this isn't something of great importance.

VI. BLANK ANALYSIS RESULTS

The blank analysis was reviewed. The contaminants in the blank are listed below:

<u>Fraction</u>	<u>Compound</u>	<u>Concentration</u>	<u>D.L.</u>
<u>VOA</u>	<u>Methylene Chloride</u>	<u>2.3</u>	<u>5</u>
<u>VOA</u>	<u>Acetone</u>	<u>3.1</u>	<u>5</u>
<u> </u>	<u> </u>	<u> </u>	<u> </u>

Remarks: The compounds are below the certified detection
limits. The EPA blind blank also was below the certified
detection limits.
The field blank contained the highest levels of
all compounds.

VII. DFIPP AND BFB PERFORMANCE RESULTS

The DFIPP performance results were reviewed and found to be within the specified criteria.

The BFB performance results were reviewed and found to be within the specified criteria.

The (DFIPP/BFB) performance result(s) was/were reviewed and the following abundances were found to fall outside the specified criteria:

<u>Compound</u>	<u>m/z</u>	<u>Required Abundance</u>	<u>Actual Abundance</u>
_____	_____	_____	_____
_____	_____	_____	_____
_____	_____	_____	_____
_____	_____	_____	_____

The (DFIPP/BFB) performance results which were found to be outside of the contractually required tuning requirements do not have an adverse technical impact on the data.

Remarks: _____

VIII. CHROMATOGRAPHY CHECKS

Type of Column: Packed Column _____ Fused Silica Capillary Column (FSCC) _____

Packed Column Chromatography Check

<u>Tailing Factors</u>	<u>Acceptance Windows</u>	<u>Actual</u>
Benzidine	Less than 3	_____
Pentachlorophenol	Less than 5	_____

FSCC Chromatography Check

50-ng benzidine detectable? Yes _____ No _____
Pentachlorophenol response factor? Yes _____ No _____

Remarks: _____

IX. STANDARDS

General shape of the total ion chromatogram:

	<u>Acids</u>	<u>Base/ Neutrals</u>	<u>Volatiles</u>	<u>Pesticides</u>
Peak Shape	_____	_____	_____A_____	_____
Interferences	_____	_____	_____	_____
Background	_____	_____	_____	_____

Area Response

4-nitrophenol	_____
2,4-dinitrophenol	_____
Pentachlorophenol	_____
Benzidine	_____
Hexachlorocyclopentadiene	_____
Nitrobenzene	_____
Isophorone	_____
Dinitrotoluenes	_____

Remarks: _____

Reviewer's Name: _____

FIS Telephone No.: _____

Commercial Telephone No.: _____

X. CALIBRATION VERIFICATION - OPTIONAL

(Improper calibration should be reported under Comments Section on the Introduction Page.)

Calibration verified at least once each 8-hour shift: Yes _____ No _____

Mean percent change less than 20% for:

		<u>% Change</u>
Base/Neutral Fraction:	Acenaphthene	_____
	1,4-dichlorobenzene	_____
	Hexachlorobutadiene	_____
	2-chloronaphthalene	_____
	N-nitrosodiphenylamine	_____
	Di-n-octylphthalate	_____
	Fluoranthene	_____
	Benzo(a)pyrene	_____
	Mean	_____
Date of: Calibration	_____	
Verification	_____	
Analysis	_____	
Acid Fraction:	P-chloro-m-cresol	_____
	2,4-dichlorophenol	_____
	2-nitrophenol	_____
	Phenol	_____
	Pentachlorophenol	_____
	Mean	_____
Date of: Calibration	_____	
Verification	_____	
Analysis	_____	
Volatile Fraction:	1,1-dichloroethylene	_____
	Chloroform	_____
	1,2-dichloropropane	_____
	Toluene	_____
	Ethylbenzene	_____
	Mean	_____
Date of: Calibration	_____	
Verification	_____	
Analysis	_____	

Instrument found to be in calibration during analysis: Yes _____ No _____

Remarks: _____

TABLE 1

SAMPLE SUMMARY, INTERSTATE/WOBURN

May 1, 1984

Station NUS	Location ERT	Sample Number	Date	Time	Medium	Organic Traffic Number	Well* Depth	Water* level	Material	Comments
Well 2D	Well IUS2A	76854	5/1/84	1140	Groundwater	A2750	89	4.68	Bedrock	
Well 2D	Well IUS2A	76855	5/1/84	1140	Groundwater	A2751	89	4.68	Bedrock	Replicate
Well 2M	Well IUS2B	76857	5/1/84	1320	Groundwater	A2753	55	3.40	Till	
Well 2S	Well IUS2C	76856	5/1/84	1150	Groundwater	A2752	20	2.78	Sand	
Well 3A	Well IUS3A	76858	5/1/84	1632	Groundwater	A2754	62	4.00	Bedrock	
Well 3B	Well IUS3B	76861	5/1/84	1750	Groundwater	A2757	45	4.60	Till	
Well 3C	Well IUS3C	76860	5/1/84	1718	Groundwater	A2756	24	4.70	Sand	
Blank	EPA Lab	76853	5/1/84	0915	Water	A2749				
Blank	Field	76859	5/1/84	1654	Water	A2755			Bedrock	Thru Pump

* Depth below ground surface. Surface elevations have not been surveyed at this time.

LEVEL I DATA VALIDATION SUMMARY-ORGANIC DATA

Date:

Subject: Review of Region I Contract Data; SMO Case No. 2660 SAS No. _____
 Site Name Interstate/Woburn Contract Lab Cambridge Analytical
 Contract No 68-01-679
 SMO Traffic Nos. A-2749 → A 2757
 Region IV SAD Nos: _____
 Low Level Med. Level High Level Soil/Sed. Water Waste

- I. Data Completeness VOA P B/N ACID PEST TCDD
 A - Acceptable - All items delivered.
 P - Provisional - Some items not essential for review are missing.
 U - Unacceptable - Some items essential for review are missing.

REMARKS: *Std. reference spectra was missing*

- II. Spectral Performance Criteria VOA (BFB) A B/N/A (DFTPP)
 A - Acceptable - All criteria met, spectra of good quality.
 P - Provisional - All criteria not met, spectra of reasonable quality; data useable for limited purposes.
 U - Unacceptable - Criteria not met, spectra of poor quality, data unuseable.

REMARKS:

Contract Lab Cambridge Analytical SMO Case No. 2660

- III. Blank Analysis: VOA P B/N ___ ACID ___ PEST ___ TCDD ___
- A - Acceptable - No contaminants above minimum detection limit, no interference with sample results.
- P - Provisional - Contaminants present but minimal interference with sample results.
- U - Unacceptable - Gross contamination, too much interference to use data for certain components or the entire fraction.

REMARKS: *some contaminants present and compounds therefore rejected - acetone + chloroform*

- IV. Surrogate Spike Results: VOA P B/N ___ ACID ___ PEST ___ TCDD ___
- Note: Sample data flagged on individual basis.

- A. Individual sample flagging criteria.
- Acceptable - All surrogate recoveries within criteria.
- Suspect - Any surrogate recoveries, outside criteria and/or recoveries of <10% substantiated as a matrix effect.
- Invalid - Any recoveries of <10% that are unsubstantiated as a matrix effect.

REMARKS *Toluene - d₈ just outside acceptable limit*

- B. Case Summary VOA P
- A - Acceptable - <10% of samples reported as suspect
- P - Provisional - >10% but <50% of samples reported as suspect
- U - Unacceptable - >50% of samples reported as suspect and/or any samples reported as invalid.

VOA	<u>15</u>	of	<u>36</u>	Samples Suspect;	<u>0</u>	Invalid
B/N	___	of	___	Samples Suspect;	___	Invalid
ACID	___	of	___	Samples Suspect;	___	Invalid
PEST	___	of	___	Samples Suspect;	___	Invalid
TCDD	___	of	___	Samples Suspect;	___	Invalid

One compound consistently ~ 5% lower than limit of 90%

Contract Lab Cambridge Analytical Case No. 2660

REMARKS:

V. Matrix Spike Results: VOA A B/N ACID PEST TCDD

Note: No action taken on Matrix Spike results alone.

A - Acceptable - <10% of compounds outside criteria

P - Provisional - >10% but <50% of compounds outside criteria.

U - Unacceptable - >50% of compounds outside criteria and/or any recoveries of <10%.

VOA	<u> 1 </u>	of	<u> 10 </u>	Compounds outside criteria; No. <10%	<u> A </u>
B/N	<u> </u>	of	<u> </u>	Compounds outside criteria; No. <10%	<u> </u>
ACID	<u> </u>	of	<u> </u>	Compounds outside criteria; No. <10%	<u> </u>
PEST	<u> </u>	of	<u> </u>	Compounds outside criteria; No. <10%	<u> </u>
TCDD	<u> </u>	of	<u> </u>	Compounds outside criteria; No. <10%	<u> </u>

REMARKS: 1,1-Dichloroethane 48% Rec.

VI. Duplicate Spike Results: VOA P B/N ACID PEST TCDD

A - Acceptable - <10% of compounds outside criteria.

P - Provisional - >10% but <50% of compounds outside criteria.

U - Unacceptable - >10% of compounds outside criteria.

VOA	<u> 1 </u>	of	<u> 5 </u>	Compounds outside criteria
B/N	<u> </u>	of	<u> </u>	Compounds outside criteria
ACID	<u> </u>	of	<u> </u>	Compounds outside criteria
PEST	<u> </u>	of	<u> </u>	Compounds outside criteria
TCDD	<u> </u>	of	<u> </u>	Compounds outside criteria

REMARKS: Chlorobenzene 15.8 RPD

Contract Lab Cambridge Analytical Case No. 2660

VIII. Blind Blank Results: VOA A B/N ACID PEST TCDD

No blank submitted Blank submitted-See Remarks

A - Acceptable - No compounds found, compounds found but identified in lab blanks; refer to lab blank criteria for action.

P - Provisional - Minor contaminants found inconsistent with Lab blank and also found in samples; use blind blank as blank for case and treat as lab blank with minor contamination.

U - Unacceptable - Gross contamination inconsistent with lab blank and also found in samples; use blind blank as blank for case and treat as lab blank with gross contamination.

REMARKS:

Acetone + Chloroform found at levels above all others - therefore rejected.

IX. Split Sample Results: VOA A B/N ACID PEST TCDD

No split sample Split sample-See Remarks / Replicate samples

A - Acceptable - At a concentration of greater than 10 times the minimum detection limit, the same compounds were identified in the two samples with minor differences in concentration.

P - Provisional - At a concentration of greater than 10 times the minimum detection limit, the same compounds were identified in the two samples with major differences in concentration. These discrepancies could cause the data to be useful only for limited purposes.

U - Unacceptable - At a concentration of greater than 10 times the minimum detection limit, differences were found in compound identifications in the two samples. These discrepancies could cause the results for this fraction to be used for limited purposes or unusable.

Too low for use in precise calculations

Contract Lab CAA Case No. 2660

REMARKS:

- X. Holding Times: VOA A B/N ACID PEST TCDD
- A - Acceptable - All holding times within specified limits.
- P - Provisional - Holding times not met in some cases, data useable for limited purposes.
- U - Unacceptable - Holding times not met. The data unuseable or used for limited purposes.

REMARKS: